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Result
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Abr31559 Human can	Abr31169 Human can	Abr31063 Human can	Abr30941 Human can	Abr31170 Human can	Abr30613 Human can	Abr31028 Human can	Abr30612 Human can	Abr30473 Human can	Abr31271 Human can	Abr30940 Human can	Abr30509 Human can	Add23419 Breast ca	Add23767 Breast ca	Aae34759 Streptomy	Aec01228 SARS vira	Ady49658 Severe ac	Aby01491 SARS coro	Abr04849 Human can	Adz50925 Y. pestis	Aby01546 SARS coro

ALIGNMENTS

Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.

Minor histocompatibility antigen HA-1 T-cell epitope

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20-MAY-1999 AAW99195; AAW99195

(first entry)

standard; peptide; 9 AA.

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PRESULT 1
ANAW9912
ANAW99
AC ANAW9
AC MANAW
AC MANAW
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AC A
The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                                                                                                                                                                                                                             A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
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ADG66271 ADG66270

Aaw64577 Adg66269 Adg66272 Adg66271

Human pro Phosphoen

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RESULT 2
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ID AAW9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A new minor histocompatibility antigen, HA-1 - useful to treat immun diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.
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AAW99197 standard; peptide; 9
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                                                                                                                                                                                                                                              T-cell epitope; minor histocompatibility antigen transplant rejection; Graft-versus-Host Disease; neoplastic haematopoietic cell.
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                                                                       Misc-difference
                                                                                                                                                                                  Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW97572;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                         from the minor histocompatibility antigen HA-1.
                                                                                                      Location/Qualifiers
                                     /note= "His or Arg"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DF,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Engelhard
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ¥;
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2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HA-1 - useful to treat immune versus graft disease in bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                            HA-1; vaccine; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases.

Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B-and T-cells
Typing minor histocompatibility antigen HA-1 -
                                                 WPI; 1999-142960/12
                                                                                                  Goulmy
                                                                                                                                                                                                23-JUL-1997;
02-JUN-1998;
                                                                                                                                                                                                                                                                            23-JUL-1998;
                                                                                                                                                                                                                                                                                                                             04-FEB-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HA-1 H-allele sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-142855/12
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                                                                                                                                                    (UYLE-) RIJKSUNIV LEIDEN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immunogenic peptide from minor histocompatibility
for inducing tolerance to transplants and prevent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
les 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 39; 57pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ĀĀ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease.
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98EP-00870125.
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2e+06;
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by amplifying
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RESULT 6
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The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used
                                                                                                                                                                                                        Typing minor histocompatibility antigen HA-1 - identifying HA-1 R and HA-1 H alleles of HA-1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                             Claim 13; Fig
                                                                                                                                                                                                                                                           WPI; 1999-142960/12.
                                                                                                                                                                                                                                                                                                                                                      23-JUL-1997;
02-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                      04-FEB-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HA-1 R-allele
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               identifying HA-1 R and HA-1 H alleles of HA-1, used for, of genetic aberrances.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW97374
                                                                                                                                                                                                                                                                                                                       (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9905313-A2
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                                                                                                                                                           5; 59pp;
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98EP-00870125.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
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88.9%;
                                                                                                                                                             English.
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2e+06;
                                                                                                                                                                                                          by amplifying and used for, e.g. det
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                                                                                                                                                                                                           detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 7
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                                                                                               The invention relates to a novel method for providing epitopes of allelic convariants of antigenic proteins from specific species based on single concleotide polymorphism (SNP), by defining target protein/peptide or its combet, screening database of DNA encoding target protein identifying, considered by DNA sequence having SNP, creating variant epitopes, considered by DNA sequence having SNP, creating variant epitopes, considered by DNA sequence having SNP, creating variant epitopes, considered by DNA sequence having SNP, creating variant epitopes, considered by DNA sequence having SNP, creating variant epitopes, considered by DNA sequence having SNP, creating variant epitopes, considered by DNA sequence in a vaccine. The method is considered by applying the method for several protein from the considered by applying the method for several protein from the considered by applying the method for several protein from the considered by applying the method for several protein from the considered polymorphic profile. A method of the invention considered for transplanting hatcomer, preferably leukaemia, and for determining concert preferably leukaemia, and for determining concert preferably leukaemia, and for determining concert in the exemplification of the invention.
 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Strittmatter W, Moll H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUN-2002; 2002EP-00013423
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human; cytostatic; vaccine; SNP profile; cancer; leukaemia;
minor histocomptibility antigen; mHAg; T cell epitope.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 82; 119pp; English.
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                                                                    Sequence 9
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Local Similarity nes 8; Conserv
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8; Conserv
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                   97.3%;
88.9%;
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Pred. No. 2e+0
0; Mismatches
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                   Score 36; DB 8;
Pred. No. 2e+06;
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2e+06;
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RESULT 8
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                                                                                                                                                                    The invention relates to a novel method for providing epitopes of allelic covariants of antigenic proteins from specific species based on single concleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein/peptide or its concleoting allelic peptide/protein variants, expression product or its concleoting epitopes binding to MHC protein. A protein of the invention has covariating epitopes binding to MHC protein. A protein of the invention has covariating encivity, and may have a use in a vaccine. The method is concertain for generating a SNP profile of one or more individuals from a complex protein by applying the method for several protein from the concert file is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer in the SNP-related polymorphic profile. A method of the invention concerting the SNP-related polymorphic profile. A method of the invention concert in the present concerts and treating cancer, preferably leukaemia, and for determining concerts used in the exemplification of the invention.
                                                                    Query Match
Best Local S
Matches 8
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                                                                                                                                         Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 82; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2004-082200/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JUN-2002; 2002EP-00013423
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8; Conser
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                                                                      Score 36; DB
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AAW99199 standard; peptide; 13

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RESULT 10
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                                                      Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
           WO9905174-A1
                                                                                                       DH cell KIAA0223 protein sequence.
                                                                                                                                20-MAY-1999
                                                                                                                                                                            AAW99198 standard; peptide;
                                                                                                                                                                                                                                                                                                                                       Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 31; 47pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
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88.9%;
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Pred. No.
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Best Local S
Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency diseases. The present sequence represents a KIAA0223 sequence given in the present invention
                                                                     Goulmy EAJM,
                                                                                                                                           23-JUL-1998;
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                                                                                                                                                                                          WO9905173-A1.
                                                                                                                                                                                                                Homo sapiens.
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                                                                                             (UYLE-) RIJKSUNIV LEIDEN.
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nes 8; Conserv
                                   1999-142855/12.
DB; AAX16081.
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                                                                                                                                                                                                                                                                                     polymorphism in HA-1 positive homozygous individuals
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Pred. No. 1.8;
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Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-

versus-host disease

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RESULT 12
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune disease; neoplastic haematopoietic cell; KIAA0223 polymorphism.
The present sequence represents the KIAA0223 polymorphism in HA-1 negative homozygous individuals. The specification describes an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HIA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B- and T-cells
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                                                                                                                                                                                                Disclosure; Page 38;
                                                                                                                                                                                                                                   versus-host
                                                                                                                                                                                                                                                 Immunogenic peptide from minor histocompatibility antigen HA-1 - usef for inducing tolerance to transplants and prevent rejection or graft-
                                                                                                                                                                                                                                                                                                         N-PSDB; AAX16080.
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                                                                                                                                                                                                                                   disease
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Pred. No.
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                                                                                                          The invention relates to inhibiting or increasing programmed cell death of a cell. The method involves upregulating or inhibiting, respectively, the expression or activity of basic transcription factor (BTP) or its homologue in the cell. The BTP3 polypeptides and nucleic acids are useful for inhibiting or increasing programmed cell death. They are used for screening for an agent that increases or inhibits programmed cell death. The screened agent that increases or inhibits programmed cell death. The screened agent that increases or inhibits programmed cell death, is used for diagnosing or treating cancer or neurodegenerative diseases (e.g. amyotrophic lateral sclerosis, Alzheimer's disease, Parkinson's disease or multiple sclerosis), atherosclerosis, or rheumatoid arthritis. Sequences ABP71106-123 represent CARD regions of various CARD proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BTF3; cell death; apoptosis; basic transcription factor; cytostatic; nootropic; neuroprotective; antiparkinsonian; antiarteriosclerotic; antirheumatic; antiarthritic; gene therapy; CARD; E10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-MAY-2002; 2002WO-US016230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E10 protein CARD region fragment.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABP71119 standard; peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                        Inhibiting or increasing programmed cell death of a cell, for treating e.g. cancer, comprises upregulating or inhibiting, respectively, the expression or activity of basic transcription factor (BTF)3 or its
                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-167228/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-MAY-2001; 2001US-0292559P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200295001-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unidentified.
                                                                                   Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                          homolog in the cell.
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RESULT 15
AAW73901
ID AAW73
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CC chcention

CC polypeptide in an immune response provoking vaccine in chaption of a number of specific CTL).
                                                                                                                                                                                             Query Match
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Matches 4
                AAW73901;
                                            AAW73901 standard; peptide;
                                                                                                                                                                                                                                                            Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 16; SEQ ID NO 919; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-758334/74.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human respiratory syncytial virus; cytotoxic T lymphocyte; CTL; MHC epitope; DNA-based immunisation; RSV infection; virucidal; vaccine.
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Search completed: April 6, 2006, 16:57:02 Job time : 116 secs

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                                                                                           Matches
                                                                   Query Match
                                                                                          Sequence 10 AA;
                                                                                                              This sequence is a fragment of the Bacteroides forsythus protease of the invention. The protein has protease activity and haemolytic activity. The DNA and the polypeptide are useful for diagnosis, treatment and research of periodontosis. (Updated on 17-OCT-2003 to standardise OS field)
                                                                                                                                                                       Disclosure; Page 13; 13pp; Japanese.
                                                                                                                                                                                           New protease gene derived from Bacteroides forsythus - useful for diagnosis and treatment of periodontosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                           17-OCT-2003
12-APR-1999
                                                                                                                                                                                                                                 WPI; 1999-145900/13.
                                                                                                                                                                                                                                                                              25-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                             Protease; haemolytic activity; diagnosis; periodontosis.
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                                                                                                                                                                                                                                                        (SUNZ ) SUNSTAR CHEM IND CO LTD
                                           Local Similarity nes 4; Conserv
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OM protein -Run on: protein search, using sw model April Copyright 6, 2006, 16:57:26 ; Search time 19 Seconds (without alignments)
45.576 Million cell updates/sec GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd.

Title: Perfect score: Sequence: US-10-791-217A-1 37 VLXDDLLEA 9

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5 283416 seqs, 96216763 residues

Searched:

Total number of hits satisfying chosen parameters:

2523

Minimum DB | Maximum DB | seq length: 0 seq length: 15

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : PIR 80:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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hypothetical prote		protein kinase C i	()	light-harvesting p	T cell receptor al	GTP-binding protei	self-incompatibili	ribonuclease M (EC	photosystem I 13K	tubulin alpha-chai	argA protein - Sal	hypothetical prote	hypothetical prote	photosystem I 9K c	carnitine medium/1	hypothetical prote	hucolin, 75K chain	collagen type I -	PS I complex subun	acyl-[acyl-carrier	chlorophyll a/b-bi	m.	T-cell receptor be	۳.			unspecific monooxy	calnexin - mouse (ion	

ALIGNMENTS

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unspecific monooxygenase (EC 1.14.14.1) isozyme E, phenobarbital-inducible, hepatic - ra NyAlternate names: cytochrome P450
C;Species: Rattus norvegicus (Norway rat)
C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988 #text_change 31-Dec-2004
C;Accession: C26997
R;Graves, P.E.; Kaminsky, L.S.; Halpert, J.
Biochemistry 26, 3887-3894, 1987
A;Title: Evidence for functional and structural multiplicity of pregnenolone-16-alpha-ca A;Eeference number: A26997; MUID:88000604; PMID:3651420
A;Accession: C26997
A;Molecule type: protein
A;Residues: 1-10 <GRA>
A;Cross-references: UNIPROT:Q06884; UNIPARC:UPI0000174DA8
C;Superfamily: cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase; oxidoreductase; transmembrane RESULT 2 C26997

Matches Query Match Best Local 9 Local Similarity 4; Conservative 48.6%; Score 18; DB 2; Length 10; Pred. No. 9e+02; 0; Mismatches 1; Indels 0, Gaps 0,

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R;Ohishi, N.; Imaoka, S.; Suzuki, T.; Funae, Y. Biochim. Biophys. Acta 1188, 227-236, 1993
A;Title: Characterization of two P-450 isozymes placed in the A;Reference number: S39761; MUID:94072607; PMID:8251521
A;Accession: S39762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Rattus norvegicus (Norway rat)
C;Date: 19-May-1994 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
C;Accession: S39762
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S39762
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Biochemistry 26, 3887-3894, 1987
A;Title: Evidence for functional and structural multiplicity of pregnenolone-16-alpha-ca
A;Reference number: A26997; MUID:88000604; PMID:3651420
A;Accession: A26997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 unspecific monooxygenase (EC 1.14.14.1) cytochrome N;Alternate names: cytochrome P450b C;Species: Rattus norvegicus (Norway rat) C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988
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                                                                                                                                                                                          virg protein - Agrobacterium sp. (fragment)
C;Species: Agrobacterium sp.
C;Date: 19-Mar-1997 #sequence_revision 01-Aug-1997 #text_change 09-Jul-2004
C;Accession: S13224
R;Tamamoto, S:; Aoyama, T.; Takanami, M.; Oka, A.
J. Mol. Biol. 215, 537-547, 1990
A;Title: Binding of the regulatory protein Virg to the phased signal sequences upstream A;Reference number: S13224; MUID:91039316; PMID:2231718
A;Accession: S13224
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A;Residues: 1-12 <OHI>
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C;Superfamily: cytochrome P450 homology
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A;Molecule type: protein
A;Residus: 1-10 «TAM»
A;Cross-references: UNIPROT:Q7M0P7; UNIPARC:UPI000017A9C4
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                                                                                            chlorophyll a/b-binding protein Lhcb5 - spinach (fragment)
N;Alternate names: light-harvesting complex LHCIIc protein
C;Species: Spinacia oleracea (spinach)
C;Date: 19-Mar-1997 #sequence_revision 23-Apr-1999 #text_change
C;Accession: S50900
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S50900
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C; Keywords: T-cell receptor
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A; Residues: 1-11 <GOL>
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                                      R;Walters, R.G.; Ruban, A.V.; Horton, P.
Bur. J. Biochem. 226, 1063-1069, 1994
A;Title: Higher plant light-harvesting complexes LHCIIa
A;Reference number: S50900; MUID:95112835; PMID:7813461
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                          A;Accession: S50900
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R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B. J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental A;Reference number: PH0891; MUID:92078857; PMID:1836012
A;Accession: PH0924
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T-cell receptor beta chain V-D-J region (isolate 10) - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
C;Accession: PH9924
                                                                                                                                                                                                                  A;Title: Calcium-activated potassium channels
A;Reference number: JH0697; MUID:92360298; PMI
A;Accession: PS0443
A;Status: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Residues: 1-13 <ADE>
                                                                                                                                                                                                                                                                                                                                                                                               potassium channel protein Slo G3 - fruit fly (Drosophila melanogaster) (fragment) C;Species: Drosophila melanogaster C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Feb-1997 C;Accession: PS0443
                                                                                                                                                                                                                                                                                                                                                     R;Adelman, J.P.; Shen, K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrutta, Neuron 9, 209-216, 1992
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A;Experimental source: concanavalin A-activated lymphoblast
                                                                                       A;Cross-references: FlyBase:FBgn0003429
C;Keywords: alternative splicing; ion c
                                                                                                                                                                               C; Comment: This potassium channel is activated by calcium
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Pred. No.
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PMID:1497890
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K; Iwasaki, Y.; Ishikawa, H.; Hibino, T.; Takabe, T.
Biochim. Biophys. Acta 1059, 141-148, 1991
A;Title: Characterization of genes that encode subunits of cucumber A;Reference number: A56819; MUID:91355209; PMID:1883835
A;Recession: E56819
RESULT 11
562675
collagen type I - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 09-Jul-2004
C;Accession: S62675
C;Accession: S62675
R;Mizuno, M.; Kitafima, T.; Tomita, M.; Kuboki, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PS I complex subunit 8 - cucumber (fragment)
c;Speciles: Cucumis sativus (cucumber)
C;Date: 05-Jan-1996 #sequence_revision 05-Jan-1996 #text_change 09-Jul-2004
C;Accession: E56819
R;Iwasaki, Y.; Ishikawa, H.; Hibino, T.; Takabe, T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acyl-[acyl-carrier-protein] desaturase (EC 1.14.19.2) - avocado (fragments) C;Species: Persea americana (avocado) C;Date: 24-Jan-1992 #sequence_revision 24-Jan-1992 #text_change 03-Jun-2002 C;Accession: C39170 R;Shanklin, J.; Somerville, C. Proc. Natl. Acad. Sci. U.S.A. 88, 2510-2514, 1991 Proc. Natl. Acad. Sci. U.S.A. 88, 2510-2514, 1991 A;Title: Stearoyl-acyl-carrier-protein desaturase from higher plants is struk,Reference number: A39170; MUID:91172837; PMID:2006187 A;Accession: C39170
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C39170
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A;Cross-references: UNIPARC:UPI0000178182
C;Superfamily: chlorophyll a/b-binding protein
C;Keywords: chlorophyll; chloroplast; light-harvesting complex;
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A;Note: sequence extracted from NCBI backbone (NCBIP:5860
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A;Residues: 1-14 <SHA>
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Pred. No. 2.3e+03
1; Mismatches
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A;Status: preliminary
A;Molecule type: DNA
A;Molecule type: DNA
A;Residues: 1-8 <KAW>
A;Cross-references: UNIPROT:P95412; UNIPARC:UPI000017A96A; DDBJ:D50473; NID:g1217594
A;Note: this ORF is not annotated in GenBank entry PSENIRC, release 113.0
C;Superfamily: Pseudomonas stutzeri nirD protein
                                                                                                                                                                                                                                                                                                                Gene 167, 87-91, 1995
A;Title: Sequencing and characterization of the downstream y for biosynthesis of heme dl.
                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein 8 [imported] - Pseudomonas aeruginosa (fragment) C;Specias: Pseudomonas aeruginosa (C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 31-Dec-2004 C;Accession: PC4131 R;Kawasaki, S.; Arai, H.; Igarashi, Y.; Kodama, T.
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A;Molecule type: protein
A;Residues: 1-7 <EDG>
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C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
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A;Title: The osteoblastic MC3T3-E1 cells synthesized C-terminal propeptide of type I A;Reference number: S62675; MUID:97386332; PMID:9244181
A;Accession: S62675
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-15 <MIZ>
                                                                                                                                                                                                                                                                                     A;Reference number: JC4552; MUID:96144254; PMID:8566817 A;Accession: PC4131
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A;Title: Hucolin, a new corticosteroid-binding protein from A;Reference number: S68004; MUID:96087107; PMID:7498469
A;Accession: S68004
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carnitine medium/long chain acyltransferase (EC 2.3.1.-) - rat (fragment)
N;Alternate names: endoplasmic reticulum protein ERp61; glucose regulated protein GRP58;
C;Species: Rattus norvegicus (Norway rat)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 07-May-1999
C;Accession: A61354
R;Murthy, M.S.R.; Pande, S.V.
Mol. Cell. Biochem. 122, 133-138, 1993
A;Title: Carnitine medium/long chain acyltransferase of microsomes seems to be the previ A;Reference number: A61354; MUID:94049728; PMID:8232244
A;Accession: A61354
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A;Accession: A0154
A;Acces
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C;Species: Spinacia oleracea (spinach)
C;Apate: 19-Mar-1997 #sequence_revision 13-Mar-1998 #text_change 09-Jul-2004
C;Accession: S14316
R;Ikeuchi, M.; Inoue, Y.
FEBS Lett. 280, 332-334, 1991
A;Title: Two new components of 9 and 14 kDa from spinach photosystem I complex.
A;Accession: S14316; MUID:91192162; pMID:2013332
A;Accession: S14316
A;Accession: S14316
A;Accession: S14316
A;Cross-references: UNIPROT:Q7M1J1; UNIPARC:UPI000017AF28
A;Cross-references: UNIPROT:Q7M1J1; UNIPARC:UPI000017AF28
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J. Biol. Chem. 269:12263-12268(1994).
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                                                                                                                                                                                                                                                                                ECOLI
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3; Conserv
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3; Conserv
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                                                                                                                                                                                                                                                                                                                                                           IIEDDL
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ITDDVFQA 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                54.1%;
nilarity 50.0%;
Conservative
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             PubMed=1995588
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37.5%;
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MW=30142.85; METHOD=Electrospray.
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                                                                                                          Gammaproteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 20; DB
Pred. No. 5.2e
2; Mismatches
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No.
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RESULT 5
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Q9SDG CHRVI
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J. Bacteriol. 173:1367-1375(1991).
EMBL; M63619; AAA24556.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE-95392215; PubMed=7663175; DOI=10.1006/prep.1995.1049; De Pauw P., Neyt C., Vanderwinkel E., Wattlez R., Falmagne P. "Characterization of human serum N-acetylmuramyl-L-alanine ampurified by affinity chromatography."; Protein Expr. Purif. 6:371-378(1995).
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GO; GO:0008745; F:N-acetylmuramoyl-L-alanine amidase activity;
GO; GO:0001519; P:peptide amidation; NAS.

SEQUENCE 15 AA; 1600 MW; 9016B00FFF9E780A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q9R5D6;
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                              Bacteria; Proteobacteria; Gamm
Chromatiaceae; Allochromatium.
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                                                                         for poly(3-hydroxyalkanoic
D.";
                                                                                                                                                                                                                                                                                                 NCBI_TaxID=1049;
                                                                                                                                                                                                                                                                                                                                                                                                                    Chromatium vinosum.
      FEMS Microbiol.
SEQUENCE 15 AJ
                                                                                                     Liebergesell M., Schmidt B., Steinbuchel "Isolation and identification of granule for poly(3-hydroxyalkanoic acid) biosynth
                                                                                                                                                                                                      PROTEIN SEQUENCE.
MEDLINE=93146381; PubMed=1490603; DOI=10.1016/0378-1097(92)90031-I;
                                                                                                                                                                                                                                                                                                                                                                                                                                            (Fragment).
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      AA;
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          Lett. 78:227-232(1992).
A; 1874 MW; 165FA9A16BCA2A9D
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                                                                                                        granule-associated proteins relevant biosynthesis in Chromatium vinosum
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J. Biol. Chem. 267:12016-12021(1992)
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MEDLINE=92291078;
Subramaniam V.N.,
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurogna
Muridae; Murinae; Rattus.
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Della-Porta A.J.;
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Paramyxoviridae; Paramyxovirinae; Avulavirus.
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Q718K6 9PARA PRELIMINARY;
AC Q718K6;
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RT motifs as markers for viral li
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Carbohydrate metabolism; Direct protein sequencing; Isomeras
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Biochem. J
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P14405;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=90104230; PubMed=2604694;
Vangrysperre W., Ampe C., Kersters-Hilderson H., Tempst P.;
"Single active-site histidine in D-xylose isomerase from Streptomyces
violaceoruber. Identification by chemical derivatization and peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Streptomyces violaceoruber.
Bacteria; Actinobacteria; A
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10-MAY-2005 (Rel.
Xylose isomerase
                                                                                                                                                                                                                                                                          use
                                                                                                                                                                                                                                                                                                  This Swiss-Prot entry is copyright. It is produced through a obstween the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                        -!- SUBUNIT: Homotetramer.
-!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: Belongs to the xylose isomerase family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biochem. J. 263:195-199(1989).
-!- CATALYTIC ACTIVITY: D-xylose = D-
-!- COFACTOR: Binds 2 magnesium ions
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Avian Pathol. 32:361-373(2003).
EMBL; AF542917; AAQ11642.1; -;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROTEIN SEQUENCE.
STRAIN=LMG 7183;
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Paramyxoviridae; Paramyxovirinae; Avulavirus.
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
NADH:ubiquinone oxidoreductase (Complex I) iron-sulfur
fraction 20 kDa polypeptide peptide T-8 (Fragment).
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01-JUN-2002 (TrEMBLrel. 21,
Glycoprotein H-B N-TERMINAL,
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                  PROTEIN SEQUENCE.

MEDLINE-91340701; PubMed-1874725;

Christie D.L., Batchelor D.C., Palmer D.J.

"Identification of kex2-related proteases
partial amino acid sequence analysis.";

J. Biol. Chem. 266:15679-15683(1991).
                                                                                                                                                           Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; !
Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla;
Pecora; Bovidae; Bovinae; Bos.
NCBI_TaxID=9913;
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Bos taurus (Bovine).
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J. Blochem. 110:575-882(1991).
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Masui R., Wakabayashi S., Matsubara
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Mammalia; Eutheria; Laurasiatheria; Cetartiodactyla;
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"POL and K17 markers from Beltsville CTV isolates.";
submitted (SEP-2004) to the EMBL/GenBank/DDBJ databa
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3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
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RESULT 1 US-09-269-250E-18 Sequence 18, Application US/09269250E GENERAL INFORMATION: APPLICANT: GOULMY, E18a TITLE OF INVENTION: METHOD FOR TYPIN FILE REFERENCE: 58994 CURRENT APPLICATION NUMBER: US/09/26 CURRENT ETING DATE: 1999-05-21 NUMBER OF SEQ ID NOS: 38 SOFTWARE: Patentin version 3.1 SEQ ID NO 18 LENGTH: 9 TYPE: PRT		45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	
ULT 1 09-269-250E-18 09-269-250E-18 equence 18, Application US/ ENERAL INFORMATION: APPLICANT: Goulmy, Elsa TITLE OF INVENTION: METHOD I FILE REFERENCE: 58994 CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 1999-0 NUMBER OF SEQ ID NOS: 38 SOFTWARE: PatentIn version 3 EXECUTE: PRT TYPE: PRT						21				22										
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Patent No. 6830883
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PILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTMARE: Patentin version 3.1
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LENGTH: 9
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Conservative

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Score 36; DB 2; Pred. No. 4.6e+05; 0; Mismatches)

Length 9; Indels

0,

Gaps

0,

Fragments

S

VLXDDLLEA 9

Best Local Matches

Similarity 8; Conserv

Conservative

88.9%;

Pred. No. 4.6e+05; 0; Mismatches 1;

Indels

0,

Gaps

0

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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILLING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILLING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 9
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US-09-269-250E-29
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US-09-489-760-1
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Patent No. 6830883

GENERAL INFORMATION:
APPLICANT: Goulimy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 29
LENGTH: 9
LENGTH: 9
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                                                   Matches
                                                                Query Match
Best Local :
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LOCATION: (3)...(3)
OCHER INFORMATION: Xaa represents a histidine (H) or an arginine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
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ORGANISM: Human
                                                                                                                             FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (3)
OTHER INFORMATION: AMINO ACID X REPRESENTS A HISTIDINE OR AN ARGININE
OTHER INFORMATION: RESIDUE
                                                                                                                                                                                                                 TYPE: PRT ORGANISM: Histocompatibility antiger
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les 9; Conserv
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9; Conserva
1 VLXDDLLEA 9
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                                                   Score 36; DB; Pred. No. 4.6
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CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LENGTH: 9
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US-09-489-760-5
; Sequence 5, Ap
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US-09-489-760-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT; ORGANISM: Histocompatibility antigen US-09-489-760-2
RESULT 7
US-09-269-250E-26
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Best Local S
Matches
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                                                                                                                                                                                                                                                    SEQ ID NO 5
LENGTH: 9
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APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                Query Match
Best Local
                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                TYPE: PRT ORGANISM: KIAA0223 partial complementary DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 VLXDDLLEA 9
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88.9%;
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88.9%;
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
0; Mismatches 1
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Sequence 26, Application US/09269250E Patent No. 6830883
GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa

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                                                                                                       APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
FULRENT APPLICATION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION DATE: 1998-07-23
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US-09-269-250E-28
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Best Local S
Matches 8
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CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 28
LENGTH: 13
TYPET- TENT
                                                        SOFTWARE: PatentIn Ver. SEQ ID NO 14
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Best Local Similarity
Matches 8; Conserv
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APPLICANT: GOULTY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
                                                                                                                                                                                                                                                                                                            Patent No. 687837
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CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
                                                                                          NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 58994
                    LENGTH: 13
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
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TYPE: PRT
ORGANISM: Artificial Sequence
ORGANISM: Unknown Organism
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88.9%;
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Pred. No. 0.23;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 1000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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US-09-489-760-16
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Best Local S
Matches 8
                                                                                                                             SOFTWARE: PatentIn version 3.1
SEQ ID NO 38
                                                                                                                                                                                                                                                              Sequence 38, Application US/09269250E Patent No. 6830883
GENERAL INFORMATION:
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SEQ ID NO 16
   Matches
                                   Query Match
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APPLICANT: Rijksuniversiteit Te Leidin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                            APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
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                                                                               TYPE: PRT
ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Unknown Organism FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Unknown Organism:HA-1+/+
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                                                                                                                 ENGTH: 8
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 Similarity 5; Conserv
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88.9%;
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$; Score 23; DB:
$; Pred. No. 4.6;
0; Mismatches
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Pred. No. 0.23;
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Pred. No. 0.23;
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               DB 2; L
4.6e+05;
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RESULT 12
US-08-940-095-104
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Best Local Similarity
Watches 5; Conserva
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; MOLECULE TYPE:
US-08-940-095-104
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                                                                                                                                                      Sequence 104, Application US/08940093 Patent No. 6037323
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GENERAL INFORMATION:
                                                                                                                                      Patent No. 6037323
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO:
                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: COTUZZÍ, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 00:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Dufourcq, Jean
TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
               APPLICANT: Buttner, Klaus
APPLICANT: Cormut, Isabelle
APPLICANT: Metz, Gunther
TITLE OF INVENTION: APOLIPOE
TITLE OF INVENTION: AND THEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 29-SEP CLASSIFICATION: 514
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STATE: NY
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1155 Avenue of the Americas
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                                                                                                     Dasseux, Jean-Louis
Sekul, Renate
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Cornut, Isabelle
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29-SEP-1997
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                 APOLIPOPROTEIN A-I AGONISTS
AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
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Pred. No. 1.4e+02;
2; Mismatches 1
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Best Local Similarity
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APPLICANT: Dasseux, Jean-Louis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
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APPLICANT:
APPLICANT:
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FILING DATE: 29-SEP-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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                                                                                                   COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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                                 SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                           TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS
TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS
NUMBER OF SEQUENCES: 258
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                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
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OPERATING SYSTEM:
SOFTWARE: FastSE
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CITY: New York
STATE: NY
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STRANDEDNESS: sir
TOPOLOGY: linear
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REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 00
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TELEFAX: 650-493-5556
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STREET: 1155 Avenue of the Americas
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APPLICATION NUMBER: US/0 FILING DATE: 29-SEP-1997
                                                                 COMPUTER: IBM CON
OPERATING SYSTEM:
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STREET: 1155 Avenue of the Americas
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Cornut, Isabelle
Metz, Gunther
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                   US/08/940,096
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Pred. No. 1.4e
2; Mismatches
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CLASSIFICATION: 530 PRIOR APPLICATION NUMBER:

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Query Match
Best Local Similarity
Watches 5; Conserve
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US-09-465-719-104
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,093
FILING DATE: 29-SEP-1997
ATTORNEY AGENT INFORMATION:
NAME: COTUZZI, LAUTA A
REGISTRATION NUMBER: 009196-0006-999
REFERENCE/DOCKET NUMBER: 009196-0006-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEPHONE: 650-493-4935
TELEFAX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 104:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
CTARACTERISTO acid
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APPLICANT: Dasseux, Jean-Louis

APPLICANT: Sekul, Renate

APPLICANT: Sekul, Renate

APPLICANT: Ocrnut, Isabelle

APPLICANT: Metz, Gunther

TITLE OF INVENTION: APOLIPOPROTEIN A-I AGONISTS

TITLE OF INVENTION: AND THEIR USE TO TREAT DYSLIPIDEMIC DISORDERS

NUMBER OF SEQUENCES: 258

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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NAME: COTUZZI, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0005-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEPAX: 650-493-5556
TELEPAX: 650-493-5556
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: NY
                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/465,719 FILING DATE: CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
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; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: NO. 6265377e
US-09-465-719-104

Query Match
Best Local Similarity 62.2%; Score 23; DB 2; Length 15;
Best Local Similarity 62.5%; Pred. No. 1.4e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps

Qy 2 LENDLLEA 9
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Db 2 LINELLEA 9
Search completed: April 6, 2006, 17:05:27
Job time: 27.6667 secs
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Title:
Perfect score:
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No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Match
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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37
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    Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
      GenCore version (c) 1993 - 2006
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US-10-623-176-1
US-10-623-176-2
US-10-623-176-10
US-10-791-217-2
US-10-791-217-2
US-10-791-217-3
US-10-791-217-40-18
US-11-007-740-20
US-11-007-740-20
US-11-007-740-24
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US-10-791-217-14
US-10-791-217-14
US-10-623-176-42
US-11-007-740-28
US-11-007-740-28
US-11-623-176-42
US-10-623-176-42
US-10-623-176-43
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Biocceleration Ltd.
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11, 7
118, 7
20, 7
24, 7
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26,
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41,
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RESULT 2 US-10-623-176-2

Sequence 2, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.

A WRY	US-	· · · · · · · · · · · · · · · · · · ·	RES US- ; S ; G		
Query Match Best Local Matches	TYPE: PRI TYPE: PRI ORGANISM: FEATURE: OTHER INF OTHER INF OTHER INF FEATURE: NAME/KEY: LOCATION: 10-623-176		RESULT 1 Sequence 1, Application US/1062317; Sequence 1, Application US/1062317; Publication No. US20040092446A1; GENERAL INFORMATION: APPLICANT: Goulmy, Els A.J.M.; APPLICANT: Hunt, Donald F.; APPLICANT: Engelhard, Victor H.; TITLE OF INVENTION: HA-1 epitopes; FILE REFERENCE: 2183-6047US	ງ ເວລ ເປັດ α α α α α α α α α α α α α α α α α α α	31 32 32 33 34 35 35 36 36 37 37 37 37 37 37 37 37 37 37 37 37 37
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imilarity ; Conservat VLXDDLLEA 9	al De wh	TT APPLICATION NUMBER: US/10/623,17 TF FILING DATE: 2003-07-18 APPLICATION NUMBER: 09/489,760 FILING DATE: 2000-01-21 APPLICATION NUMBER: BP 97202303.0 FILING DATE: 1997-07-23 APPLICATION NUMBER: PCT/NL98/00424 FILING DATE: 1998-07-23 APPLICATION NUMBER: JP 2000-504165 FILING DATE: 2000-01-24 RE: PatentIn Ver. 2.1 RO: 0P SEQ ID NOS: 77 RE: PatentIn Ver. 2.1	ion 0400 Els Els onal rd,	N N W W M M M M M M M	
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Best Local Similarity
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PRIOR FILING DATE: 1997-07-23
PRIOR PPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR PPLICATION NUMBER: JP 2000-504165
PRIOR PILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
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; LOCATION: (1)..(9)
US-10-623-176-2
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S
                 Query Match
Best Local Similarity
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Publication No. US20040092446A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Ver. SEQ ID NO 10
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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TITLE OF INVENTION: HA-1 epitopes and uses thereof
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
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NUMBER OF SEQ ID NOS: 77
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PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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NAME/KEY: SITE
LOCATION: (1)..(9)
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                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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1 VLXDDLLEA 9
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                                       Conservative
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Pred. No. 1.7e+06;
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Pred. No. 1.7e+06;
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                                         Mismatches
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US-10-791-217-2
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                                                                                              , OTHER INFORMATION: obtained from histocompatibility antigen US-10-791-217-2
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Publication No. US20040191268A1
GENERAL INFORMATION:
                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 2
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97.3%; Score 36; DB 4; Length 9; Best Local Similarity 88.9%; Pred. No. 1.7e+06; Matches 8; Conservative 0: Migmatches
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CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
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APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
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CURRENT FILING DATE: 2004-03-02
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NAME/KBY: MISC_FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa is HISTIDINE OR ARGININE RESIDUE
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                                                                                                                                            ORGANISM: Artificial sequence FEATURE:
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100.0%; Pred. No. 1.7e+06;
tive 0; Mismatches 0;
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US-10-861-335-1
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US-10-791-217-5
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APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR APPLICATION NUMBER: DS/09/489,760
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M.
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LENGTH: 9
TYPE: PRT
ORGANISM: Artificial sequence
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                                                                                                          Query Match
                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1 SEQ ID NO 1
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Best Local
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PRIOR TILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
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88.9%;
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1.7e+06;
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RESULT 10
US-11-007-740-29
; Sequence 29, Application US/11007740
; Publication No. US20050233350A1
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Sequence 20, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
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US-11-007-740-18
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Matches 8
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Publication No. US20050233350A1

GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa

TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1

FILE REFERENCE: 2799/58994-A

CURRENT FILING DATE: 10904-12-08

PRIOR APPLICATION NUMBER: 09/169,250

PRIOR APPLICATION UMBER: 09/269,250

PRIOR PILING DATE: 1999-05-21

NUMBER OF SEQ ID NOS: 40

CONTENSED: Description of the content of the co
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PatentIn version 3.1
SEQ ID NO 20
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Best Local Similarity
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ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Exon
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Pred. No.
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Pred. No. 1.7e+06;
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PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR PILING DATE: 1997-07-23
PRIOR PELLOATION NUMBER: BCT/NL98/00424
PRIOR TILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 24
LENGTH: 10
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US-10-623-176-24
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                                                                                        RESULT 12
US-10-623-176-43
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                                 Sequence 43, Application US/10623176 Publication No. US20040092446A1 GENERAL INFORMATION:
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Best Local :
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Best Local 9
                                                                                                                                                                                                                                                Matches
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
APPLICANT: Goulmy, Els A.J.M. APPLICANT: Hunt, Donald F.
                                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: SITE
LOCATION: (1)..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-01-21
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TYPE: PRT
ORGANISM: Human
FEATURE:
HAME/KEY: MISC_FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a histidine (H) or an arginine
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide OTHER INFORMATION: wherein X can be H or R
                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                            Local Similarity
nes 9; Conserv
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hes 0;
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APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: EPT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER: DF 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER: DF 2000-504165
                                                                                                                                 ; NAME/KEY: SITE
; LOCATION: (1)..(10)
US-10-623-176-44
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US-10-623-176-44
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; NAME/KEY: SITE
; LOCATION: (1)..(10)
US-10-623-176-43
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION UNMERE: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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SEQ ID NO 44
LENGTH: 10
TYPE: PRT
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                                                                  Query Match
Best Local Similarity
                                              Matches
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TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                 FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE:
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    1 VLXDDLLEA 9
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8; Conservat
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                                              Conservative
                                                                  97.3%;
88.9%;
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Pred. No. 1.8;
                                                                       Score 36; DB
Pred. No. 1.8;
                                                Mismatches
                                                                                         DB 4;
                                                1; Indels
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                                                  Gaps
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RESULT 14 US-10-623-176-65

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APPLICANT: GOLINY, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
CURRENT FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: P07/N198/00424
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/N198/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOPTWARE: Patentin Ver. 2.1
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: DCT/NL98/00424
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 65
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US-10-623-176-74
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Publication No. US20040092446A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 8; Conserv
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Publication No. US20040092446A1
GENERAL INFORMATION:
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-10-623-176-65
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE: FRATURE: NAME/KEY: SITE LOCATION: (1)...(12)
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TYPE PT
ORGANISM: Artificial Sequence
PEATURE:
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Pred. No. 2
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; NAME/KEY: SITE
; LOCATION: (1)..(13)
US-10-623-176-74
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Search completed: April Job time: 97.6667 secs
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                                                                                                                 Matches
                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: KIAA0223
OTHER INFORMATION: sequence derived from a presumed HA-1 negative
OTHER INFORMATION: individual
FEATURE:
                                                                         1 VLXDDLLEA 9
                                                                                                                 8; Conservative
                                                       3 VIRDDLLEA 11
           6,
                                                                                                                             97.3%;
88.9%;
              2006, 17:28:45
                                                                                                                             Score 36; 1
Pred. No. 2
                                                                                                                 Mismatches
                                                                                                                                         DB 4;
                                                                                                              1; Indels
                                                                                                                                           Length 13;
                                                                                                              0,
                                                                                                                Gaps
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Minimum DB Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Result
No.
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    seq length: 0 seq length: 15
            Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*

2: /SIDSS/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/USO1_NEW_PUB.pep:*

7: /SIDSS/ptodata/2/pubpaa/USO1_NEW_PUB.pep:*

8: /SIDSS/ptodata/2/pubpaa/USO1_NEW_PUB.pep:*
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Gapop 10.0 , Gapext 0.5
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37
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          VLXDDLLEA 9
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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                                  1599
1100
1100
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                     US-11-010-748A-11
US-11-1010-748A-12
US-11-172-740-1404
US-10-927-435-63
US-10-927-634-63
US-10-927-634-63
US-10-927-634-63
US-11-129-143-169
US-11-129-143-172
US-11-129-143-172
US-11-129-143-172
US-11-129-143-173
US-11-129-143-173
US-11-021-305-62
US-11-021-305-62
US-11-021-305-65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUMMARIES
Sequence 11, Appl
Sequence 12, Appl
Sequence 1404, Appl
Sequence 63, Appl
Sequence 71, Appl
Sequence 71, Appl
Sequence 170, App
Sequence 170, App
Sequence 171, App
Sequence 172, App
Sequence 173, App
Sequence 173, App
Sequence 173, App
Sequence 65, Appl
Sequence 65, Appl
Sequence 159, App
Sequence 159, App
Sequence 159, App
Sequence 100, App
Sequence 170, App
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Description
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Qy 1 VLXDDLLEA 9 Db 1 VLHDDLEA 9	Query Match 97.3%; Score 36; DB 7; Length 9; Best Local Similarity 88.9%; Pred. No. 1.4e+05; Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;	CURRENT APPLICATION NUMBER: US/11/010,748A CURRENT FILING DATE: 2004-12-13 PRIOR APPLICATION NUMBER: PCT/EP03/06251 PRIOR PILING DATE: 2003-06-13 PRIOR FILING DATE: 2003-06-13 REGION FILING DATE: 2003-06-13 NUMBER OF SEQ ID NOS: 926 SOFTWARE: Patentin version 3.1 SEQ ID NO 11 SEQ ID NO 11 COGNAUSM: Homo sapiens PATURE: PRT OGGANUSM: Homo sapiens PATURE: NAME/KEY: MISC_FEATURE LOCATION: (1)(9) OTHER INFORMATION: ell epitopes US-11-010-748A-11	ESULT 1 IS-11-010-748A-11 S-quence 11, Application US/110107 Sequence 11, Application US/110107 Publication No. US20050244421A1 GENERAL INFORMATION: APPLICANT: Merck Patent GmbH APPLICANT: MORLTHER, Wolfgan APPLICANT: MORLTHER, Wolfgan APPLICANT: SCHARM, Burkhard TITLE DEFEDENCE. LAPPLICANTIONS.	.4 15 6 US-10-522-297-12 Sequence 12, ALIGNMENTS	\$1.4 7 US-11-225-686-3916 \$1.4 7 US-11-225-686-3932 \$1.4 7 US-11-225-686-3941 \$1.4 7 US-11-225-686-3941 \$1.4 7 US-11-202-009-2954 \$1.4 7 US-11-202-009-3916 \$1.4 9 US-10-202-009-3941 \$1.4 12 7 US-11-203-009-1739 \$1.4 12 7 US-11-203-009-1739 \$1.4 13 7 US-11-207-078-73	19 51.4 7 7 US-11-225-686-3915 Sequence 3915,
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US-11-010-748A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-11-172-740-1404
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LENGTH: 9
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1404, Application US/11172740 Pullication No. US20060057724A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: PCT/EP03/06251
PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: MASCIA, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: COMPOSITIONS AND METHODS FILE REFERENCE: MER-136
CURRENT APPLICATION NUMBER: US/11/010,748A
CURRENT FILING DATE: 2004-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: ALEXANDROV, Nickolai
APPLICANT: BROVER, Vyacheslav
TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
TITLE OF INVENTION: PLANT CHARACTERISTICS AND PHENOTYPES
FILE REFERENCE: 2750-1602PUS2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/584,829
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/584,800
PRIOR FILING DATE: 2004-06-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/583,621
PRIOR FILING DATE: 2004-06-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/11/172,740
CURRENT FILING DATE: 2005-06-30
                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 2523
                                                                                                                                                                                                                                                                                       LOCATION: (208)..(258)
OTHER INFORMATION: Pfam Name:
NAME/KEY: misc_feature
                                                                                                     OTHER INFORMATION: Utility: FEATURE:
                                                                                                                                                                                                       LOCATION: (208)..(258)
OTHER INFORMATION: Pfam
                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Arabidopsis thaliana
                         FEATURE:
                                        LOCATION:
OTHER INFORMATION: Utility: Useful for making larger plants
                                                                                                                                         LOCATION
                                                                                                                                                              NAME/KEY: misc_feature
                                                                                                                                                                                                                            NAME/KEY: misc feature LOCATION: (208)..(258)
                                                                                                                                                                                                                                                                      FEATURE:
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                                                                              NAME/KEY: misc_feature
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8; Conserv
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88.9%;
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                                                                                                                          Useful
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Pred. No. 1.4e+05;
                                                                                                                                                                                                           PHD;
                                                                                                                                                                                                                                                                                           PHD;
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                                                                                                                                                                                                           Pfam
                                                                                                                          for increasing
                                                                                                                                                                                                                                                                                           Pfam
                                                                                                                                                                                                                                                                                           Description: PHD-finger
                                                                                                                                                                                                           Description: PHD-finger
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                                                                                                                          chlorophyll and photosynthetic cap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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RESULT 4
US-10-27-435-63
J Sequence 63, Application US/10927435
Publication No. US20060045884A1
GENERAL INFORMATION:
GENERAL INFORMATION:
JEFFREY
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                                                                                               APPLICANT: MOLLDREM, JEFFREY
TITLE OF INVENTION: ANTI-CANCER VACCINES
FILE REFERENCE: UTSC:788USC1
CURRENT APPLICATION NUMBER: US/10/927,634
CURRENT FILING DATE: 2004-08-26
PRIOR FILING DATE: 2004-08-26
PRIOR FILING DATE: 2003-08-26
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PATENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                          US-10-927-634-63
                                                                                                                                                                                                                                                                                                                               RESULT 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 63
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                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                      Sequence 63, Application US/10927634
Publication No. US20060045881A1
                                                                                       SEQ ID NO 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: MOLLDREM, JEFFREY
TITLE OF INVENTION: VACCINES FOR AUTOIMMUNE AND INFECTIOUS DISEASE
FILE REFERENCE: UTSC: 788USC2
CURRENT APPLICATION NUMBER: US/10/927,435
CURRENT FILING DATE: 2004-08-26
PRIOR APPLICATION NUMBER: 60/489,238
PRIOR APPLICATION NUMBER: 60/489,238
PRIOR PLING DATE: 2003-08-26
PRIOR PILING DATE: 2003-08-26
PRIOR PILING DATE: 2003-08-26
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NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION:
OTHER INFORMATION:
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OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                   ORGANISM: Artificial Sequence
                                                     TYPE: PRT
                     FEATURE:
                                                                     ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Description of Artificial Sequence: Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      1;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for making ornamental plants with modified flowers
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; TOPOLOGY: linear; MOLECULE TYPE: peptide; SEQUENCE DESCRIPTION: SEQ ID NO: 71: US-10-353-783-71
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Best Local Similarity
Matches 5; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Peptide US-10-927-634-63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-353-783-71
; Sequence 71, Application US/10353783
; Publication No. US20050261175A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 6
                                                                                                                                  TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 71:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/353,783 FILING DATE: 28-Jan-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Suggs, Sidney V.

Martin, Francis H.

TITLE OF INVENTION: Stem Cell Factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Zsebo, Krisztina M.
Bosselman, Robert A.
                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 VLXDDLLE 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A APPLICATION DATA:
APPLICATION NUMBER: 08/448,729
FILING DATE: 24-MAY-1995
APPLICATION NUMBER: 08/172,329
FILING DATE: 21-DEC-1993
                                                                         STRANDEDNESS: single
                                                                                             TYPE: amino acid
                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 01017/32958A
                                                                                                                                                                                                                                                                      NAME: Clough, David W. REGISTRATION NUMBER: 36,107
                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/537,198
FILING DATE: 11-UUN-1990
APPLICATION NUMBER: 07/422,383
FILING DATE: 16-OCT-1989
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 07/589,701
FILING DATE: 01-OCT-1990
APPLICATION NUMBER: 07/573,616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: United States of America ZIP: 60606-6402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Chicago
STATE: Illinois
                                                                                                                                                                                               TELEFAX: 312/474-0448
                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 24-AUG-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 07/684,535 FILING DATE: 10-APR-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 07/982,255 FILING DATE: 25-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Marshall, O'Toole, STREET: 6300 Sears Tower, 233
                                                                                                                  LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  56.8%;
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Pred. No.
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South Wacker Drive
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APPLICANT: YELISEEV, Alexei A.

TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION

FILE REFERENCE: C38435/121966

CURRENT APPLICATION NUMBER: US/11/129,143

CURRENT FILING DATE: 2005-05-13

NUMBER OF SEQ ID NOS: 197

SEQ ID NO 168

LENGTH: F6
                                                                                                                                                                                                                                                                                                                                                                                    US-11-129-143-169
; Sequence 169, Application US/11129143
; Publication No. US20050266518A1
; GENERAL INFORMATION:
                                                                                                                       ; TYPE: PRT; ORGANISM: Rhizobium US-11-129-143-169
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US-11-129-143-168
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                                                             Matches
                                                                          Query Match
Best Local :
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Best Local Similarity 50.0
Matches ... 4; Conservative
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                                                                                                                                                                                                                FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                            APPLICANT: LOPEZ-ULÍBARRI, RUAL
APPLICANT: MAYER, Anne F.
APPLICANT: YELLSEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
                                                                                                                                                                                                                                                                                                                           APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, W
APPLICANT: HUMBELIN, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Bradyrhizobium japonicum
                                                                                                                                                                      ENGTH: 5
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1 DDILD 5
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                              4 DDLLE 8
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HUMBELIN, Markus
LOPEZ-ULIBARRI, Rual
MAYER, Anne R
                                                                          Similarity
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b. US20050266518A1
                                                             Conservative
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                                                                        54.1%;
                                                          Score 20; DB 7; Le
Pred. No. 1.4e+05;
2; Mismatches 0;
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Pred. No. 1.4e+05;
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Pred. No. 2.1e+02;
                                                                                       Length 5;
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US-11-129-143-172
(US-11-129-143-172); Sequence 172, Application US/11129143; Publication No. US20050266518A1; GENERAL INFORMATION:
APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner; APPLICANT: HUMBELIN, Markus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: BRETŽEL, Werner
APPLICANT: HUMBELIN, MARKUS
APPLICANT: HUMBELIN, MARKUS
APPLICANT: MAYER, Anne F.
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT APPLICATION NUMBER: US/11/129,143
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 170
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; TYPE: PRT
; ORGANISM: Bacillus stearothermophilus
US-11-129-143-170
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APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELLIN, Markus
APPLICANT: HUMBELLIN, Markus
APPLICANT: HOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
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                                                                                                                                         RESULT 11
                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Bacillus subtilis US-11-129-143-171
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                                                                                                                                                                                                                                                                                                                                                                                         SOPTWARE: PatentIn version 3.1
SEQ ID NO 171
LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 171, A Publication No.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
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3; Conserv
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lo. US20050266518A1
                                                                                                                                                                                                                                                                  Conservative
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60.0%;
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Pred. No. 1.4e+05;
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Pred. No.
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                                                                                                                                                                                                                                                                                      1.4e+05;
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TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 172
LENGTH: 5
TYPE: PRT
ORGANISM: Escherichia coli
US-11-129-143-172
                                                                                                                                                                                                                                                                  RESULT 13
US-10-982-440-126
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Matches
                                                                                                                                                                                                              Sequence 126, Application US/10982440 Publication No. US20060018909A1 GENERAL INFORMATION:
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SEQ ID NO 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 173, Appublication No.
SOFTWARE: PatentIn version 3.3
SEQ ID NO 126
LENGTH: 7
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Best Local &
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                   APPLICANT: Oliner, John
APPLICANT: Graham, Kevin
TITLE OF INVENTION: Angiopoietin-2 Specific Binding Agents
FILE REFERENCE: 04-881-A
CURRENT APPLICATION NUMBER: US/10/982,440
CURRENT FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/620,161
PRIOR PILING DATE: 2004-10-19
NUMBER OF SEQ ID NOS: 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: YELISERY, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Haemophilus influenzae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  YELISEEV, Alexei A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/11129143
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Pred. No. 1.4e+05;
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RESULT 15

US-11-021-305-65.

US-11-021-305-65.

Sequence 65, Application US/11021305

Publication No. US20050282733A1

GENERAL INFORMATION:

APPLICANT: Prins, Johannes B

APPLICANT: Hutley, Louise J

TITLE OF INVENTION: Differentiation-modulating agents and uses therefor FILE REFERENCE: DAVI69.001CP1

CURRENT FILING DATE: 2004-12-23

CURRENT FILING DATE: 2004-12-23

PRIOR APPLICATION NUMBER: US/11/021,305

CURRENT FILING DATE: 2002-06-27

NUMBER OF SEQ ID NOS: 170

SOFTWARE: PATENTIN VERSION 3.3

SEQ ID NO 65

LENGTH: 8

TYPE: PRT

ORGANISM: mammalian

US-11-021-305-65
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Publication No. US20050282733A1

GENERAL INFORMATION:

APPLICANT: Prins, Johannes B

APPLICANT: Hutley, Louise J

FILE OF INVENTION: Differentiation-modulating agents and uses therefor FILE REFERENCE: DAVI69.001CP1

CURRENT APPLICATION NUMBER: US/11/021,305

CURRENT FILING DATE: 2004-12-23

PRIOR APPLICATION NUMBER: USSN 60/392,130

PRIOR PILING DATE: 2002-06-27

NUMBER OF SEQ ID NOS: 170

SOFTWARE: PatentIn version 3.3

SEQ ID NO 62

LENGTH: 8

TYPE: PRT

COGANISM: mammalian
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; ORGANISM: Homo sapiens
US-10-982-440-126
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US-11-021-305-62
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                                                                           Query Match 54.1%; Score 20; DB 7; Les Best Local Similarity 100.0%; Pred. No. 1.4e+05; Matches 4; Conservative 0; Mismatches 0;
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SUMMARIES

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Adp26019 Plasmodiu	Adk05295 Hepatitis	-	Adq10530 Cercopith	Adk05293 Hepatitis	Adk05291 Hepatitis	Epi	MHC cla	Aay10122 T cell ep		Aaw97373 Peptide e	BCZ4 tu	Aaw97572 T-cell ep	Aaw99195 Minor his	Aaw97414 KIAA0223	Aaw99198 DH cell K	Adh40334 Human min	Aaw97374 HA-1 R-al	Aaw99197 Minor his	Aaw97415 KIAA0223	Aaw99199 vR cell K	Adh40333 Human min	Aaw97375 HA-1 H-al	inor hi	Description	

Sequence 9

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Abp71119 E10 prote Adw30865 HLA bindi	Aaw73901 B. forsyt	Adu99636 BFA5 tumo	Adt49074 Human BFA	Adt73023 Human RSV	Adm96258 Human ser	Adm96261 Human ser	Aae31275 Human mag	Adm96262 Human ser	Ady63770 Human alb	Abu97231 Enzyme po		Abu97232 Enzyme pe	Abu97233 Enzyme pe	Aab85066 Human ser	Aab85071 Human ser	Adq26722 Topo V Hr	Adt02787 Human tum	Abj20115 MHC bindi	Add23269 Breast ca

ALIGNMENTS

Minor histocompatibility antigen HA-1 T-cell epitope #2

20-MAY-1999

(first entry)

AAW99196;

AAW99196 standard; peptide; 9 AA.

RESULT 1
AAW99196
ID AAW9
XX AAW
AC AAW9
XX Mino
XX Homc
XX The
PT disk
PT dis The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease. diseases A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation. treatment of severe aplastic anaemia, leukaemia, and immune deficiency Claim 3; Page 32; 47pp; English. WPI; 1999-153312/13. Goulmy EAJM, Hunt DF, 04-FEB-1999. Homo sapiens. 23-JUL-1997; 23-JUL-1998; WO9905174-A1. (UYLE-) RIJKSUNIV LEIDEN. 97EP-00202303. 98WO-NL000425. Engelhard VH;

DB 2;

Length 9;

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RRESULT 2
AAW9737
ID AAW99
XX AAW9
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R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant;
severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. defor genetic aberrances.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-JUL-1997;
02-JUN-1998;
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ADH,40333;
                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-142960/12
                                                                                                                                                                                                                                                                                                                                                                           aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 allels. The HA-1 peptides can be used
                                              ADH40333 standard; peptide; 9 AA
                                                                                                                                                                                                                                                                                                                  Sequence
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98EP-00870125.
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Pred. No. 2e+06;
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Pred. No. 2e+06;
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RESULT 4
AAW99199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-082200/08.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-JUN-2002; 2002EP-00013423
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Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
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                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 9 AA;
                                                                                                                                           AAW99199;
                                                                                                                                                                             AAW99199 standard; peptide;
                                                                       vR cell KIAA0223 protein sequence
                                                                                                         20-MAY-1999
                                                                                                                                                                                                                                                                                                                                                     Local
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100.0%; Pred. No. 2e-
tive 0; Mismatches
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RESULT 5
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency diseases. The present sequence represents a KIAA0223 sequence given in the present inventor.
                                                                                                                                                                T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune dis neoplastic haematopoietic cell; KIAA0223 polymorphism.
         Goulmy EAJM,
                                                     23-JUL-1997;
                                                                                                                       W09995173-A1
                                                                                                                                                                                                                 KIAA0223
                                                                                                                                                                                                                                        20-MAY-1999
                                                                                                                                                                                                                                                                                  AAW97415 standard; protein; 13 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
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                                                                                                 04-FEB-1999.
                                                                                                                                           Homo sapiens.
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                                (UYLE-) RIJKSUNIV LEIDEN.
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                                                                            1998;
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         Hunt DF,
                                                     97EP-00202303
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        Engelhard VH;
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Pred. No.
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                                                                                                                                                                              autoimmune disease;
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Best Local (
The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                              A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents the KIAA0223 polymorphism in HA-1 positive homozygous individuals. The specification describes an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B- and T-cells
                                                                                                                    Disclosure; Page 15;
                                                                                                                                                                                                                   WPI; 1999-153312/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Minor histocompatibility antigen HA-1 T-cell epitope #3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             versus-host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-
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                                                                                                                   47pp; English.
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Best Local
                                                                                                      Matches
                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                      The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
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02-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. de
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98EP-00870125.
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RESULT 9
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AAW99198;

AAW99198 standard; peptide;

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20-MAY-1999

(first entry)

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                                                                                                                                                               Query Match
Best Local
                                                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-082200/08
                                                                                                                                                                                                                                                                      Sequence 9 AA;
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                                                                   1 VLHDDLLEA 9
                                                                                                                                   Similarity
8; Conserv
VLRDDLLEA
                                                                                                                                      Conservative
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                                                                                                                                                                      Score 37;
Pred. No.
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                                                                                                                                          Mismatches
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2e+06;
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Best Local
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                                                                                                                               T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune disneoplastic haematopoietic cell; KIAA0223 polymorphism.
                                                                                                                                                                                                              20-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 31; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           marrow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
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 23-JUL-1997;
                         23-JUL-1998;
                                                                                                      Homo sapiens
                                                                                                                                                                                  KIAA0223 polymorphism in HA-1 negative homozygous individuals.
                                                                                                                                                                                                                                                               AAW97414 standard;
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                                                                             WO9905173-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell KIAA0223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and organ transplantation.
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                                                                                                                                                                                                                                                                                                                                                          VLHDDLLEA 9
                                                                                                                                                                                                                                                                                                                              VLRDDLLEA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                       A,
                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                            (first entry)
97EP-00202303
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                           98WO-NL000424
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                                                                                                                                                                                                                                                               protein;
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88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                 Score 37;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                  5.1;
                                                                                                                                           Disease; autoimmune disease;
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                                                                                                                                                                                                                                                                                                                                                                                                             Length 13;
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RESULT 11
AAW99195
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents the KIAA0223 polymorphism in HA-1 negative homozygous individuals. The specification describes an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B- and T-cells
A new minor histocompatibility antigen, diseases and prevent rejection and host
                                                                                                                                                                                                                                                                                                                        Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-versus-host disease.
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N-PSDB; AAX16080.
                                                                                                                                                                             04-FEB-1999
                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                            Minor histocompatibility antigen HA-1 T-cell epitope #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW99195 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 13
                                                                   Goulmy EAJM,
                                                                                                                        23-JUL-1997;
                                                                                                                                                   23-JUL-1998;
                                                                                                                                                                                                      WO9905174-A1
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                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                              (UYLE-) RIJKSUNIV LEIDEN
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les 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                   Hunt DF,
                                                                                                                        97EP-00202303
                                                                                                                                                   98WO-NL000425
                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               82.2%;
                                                                                                                                                                                                                               His, Arg
                                                                   Engelhard
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 HA-1 - useful to treat immune versus graft disease in bone
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ARBSULT 12
ARM97572
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune disease; neoplastic haematopoietic cell.
                                                  The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides.
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Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   marrow and organ transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-142855/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-JUL-1997;
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                                                                                                                                                                                                                                                                                        Claim 1;
                                                                                                                                                                                                                                                                                                                                                versus-host
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                                                                                                                                                                                                                                                                                                                                                                     Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-
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peptides
| T-cells
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                                                                                                                                                                                                                                                                                     Page 39; 57pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                disease
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                             also
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                             be used
                                                                                                                                                                                                                                                                                     English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Engelhard VH;
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Pred. No.
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                                raise
                                antibodies,
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                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention comprises an expression vector that contains a nucleic acid encoding a tumor antigen (e.g. BCZ4 or BFY3). The expression vector of the invention is useful for the expression of multiple tumor antigens and the prevention or treatment of cancer. The present amino acid sequence represents a BCZ4 tumor antigen antigenic peptide that was used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New expression vector comprising a nucleic acid encoding a tumor antigen, e.g. BFA4, BCY1, BFA5, BCZ4, or BFY3, useful for expressing multiple tumor antigens, or for preventing or treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berinstein N,
Singh-Sandhu D;
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                      AAW97373;
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16-MAY-2003; 2003US-0471193P.
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Antibody; adipocyte; heavy chain; light chain; obesity; fat;
                       CDR region of anti-adipocyte antibody heavy chain, FAT 71.
                                                     29-AUG-2001
                                                                               AAU02706;
                                                                                                   AAU02706 standard; peptide;
                                                                                                                                                                                                                                                                            Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Typing minor histocompatibility antigen HA-1 - identifying HA-1 R and HA-1 H alleles of HA-1,
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98EP-00870125.
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2e+06;
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CC sequences of anti-adipocyte monoclonal antibody heavy chain, light chain, CC and heavy chain complementarity determining regions (CDR) of the complementarity determining regions (CDR) of the complement of the complement of the complement of the complement of complement of chair can be used in the treatment of obesity and complement yellated diseases. The antibodies can be used to deliver drugs or can be used as a therapeutic itself. Antibodies binding specifically to can be used to activate the immune system to destroy the cells by complement mediated lysis. The antibodies may be labeled with a complement mediated lysis. The antibodies may be labeled with a complement mediated lysis in human subjects e.g. to determine the presence of diagnosis in human subjects e.g. to determine the presence or level of adipocytes in a cell or tissue sample. The antibodies can be used as an alternative means of treatment for obese compatients other than undergoing surgery to remove excess fat. Antibodies compatible compatible compatible can also be produced e.g. intra-
Sequence 13 AA;
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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A;Molecule type: protein
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A;Cross-reference
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A;Title: Characterization of genomic clones specifying A;Reference number: I46868; MUID:84221901; PMID:6328491
A;Accession: I46868
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J. Biol. Chem. 267, 25703-25708, 1992
A;Title: Two isoforms of the thyrotropin-releasing hormone receptor A;Reference number: A45096; MUID:93100278; PMID:1334485
A;Accession: A45096
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C;Species: Rattus norvegicus (Norway rat)
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C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 09-Jul-2004
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A;Molecule type: mRNA
A;Residues: 1-15 <DE1>
                                                                                                                                                     R;Altmann, F.; Paschinger, K.; Dalik, T.; Vorauer, K. Eur. J. Biochem. 252, 118-123, 1998
A;Title: Characterisation of peptide-N4-(N-acetyl-bet. A;Reference number: A59272; MUID:98181894; PMID:95237A;Accession: B59272
A;Status: preliminary
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                                                                                                      R; Iwasaki, Y.; Ishikawa, H.; Hibino, T.; Takabe, T. Biochim. Biophys. Acta 1059, 141-148, 1991
A;Title: Characterization of genes that encode subunits A;Reference number: A56819; MUID:91355209; PMID:1883835
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                          A; Molecule type: protein A; Residues: 1-15 < IWA>
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GTP-binding protein o-ral - Pacific electric ray (fragment)
C;Species: Torpedo californica (Pacific electric ray)
C;Date: 22-Nov-1993 #sequence_revision 27-Feb-1997 #text_change 13-Mar-199
C;Accession: 829485
R;Volknandt, W.; Pevsner, J.; Elferink, L.A.; Scheller, R.H.
FEBS Lett. 317, 53-56, 1993
A;Title: Association of three small GTP-binding proteins with cholinergic A;Reference number: 829485; MUID:93154521; PMID:8428634
A;Accession: 829485
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: UNIPROT:Q06884; UNIPARC:UPI0000174DA8
C;Superfamily: cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase; oxidoreductase; transmembrane
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A; Accession: C26997
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Biochemistry 26, 3887-3894, 1987
A;Tille: Bridence for functional and structural multiplicity
A;Reference number: A26997; MUID:88000604; PMID:3651420
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C;Accession: C26997
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C;Species: Rattus norvegicus (Norway rat)
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C;Accession: S55696
R;Hunt, M.; Koehler, P.
Biochim. Biophys. Acta 1249, 15-22, 1995
A;Title: Purification and characterization
A;Title: Number: S55696; MUID:95284106;
virG protein - Agrobacterium sp. (fragment)
C;Species: Agrobacterium sp.
C;Date: 19-Mar-1997 #sequence_revision 01-Aug-1997 #text_change
C;Accession: S13224
R;Tamamoto, S.; Aoyama, T.; Takanami, M.; Oka, A.
J. Mol. Biol. 215, 537-547, 1990
A;Title: Binding of the regulatory protein VirG to the phased si
A;Reference number: S13224; MUID:91039316; PMID:2231718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 10
S55696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       phosphoenolpyruvate carboxykinase - Trypanosoma brucei
C;Species: Trypanosoma brucei
C;Date: 28-OCt-1995 #sequence_revision 03-Nov-1995 #text_change 09-Jul-2004
C;Accession: S55696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: UNIPROT:Q06884; UNIPARC:UPI0000174DA7
C;Superfamily: cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase; oxidoreductase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Species: Rattus norvegicus (Norway rat)
C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988
C;Accession: A26997
                                                                                                                                                                                                                                                                                                                                                                A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-9 <HUN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unspecific monooxygenase (EC 1.14.14.1) cytochrome N;Alternate names: cytochrome P450b C;Species: Rattus norvegicus (Norwav rar)
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A26997
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A;Accession: A26997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R; Graves, P.E.; Kaminsky, L.S.; Halpert,
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Pred. No. 2.6e+03
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Pred. No.
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cytochrome P450 UT-7b - rat (fragment) (;Species: Rattus norvegicus (Norway ra C;Date: 19-May-1994 #sequence_revision C;Accession: S39762 R;Ohishi, N.; Imaoka, S.; Suzuki, T.; F
                                                                                              RESULT
S39762
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PH0924
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Bur. J. Biochem. 213, 591-597, 1993
A;Title: Characterization of a clotting protein, isolated
A;Reference number: S30348; MUID:93238739; PMID:8097463
A;Accession: S30348
                                                                                                                                                                                                                                                                                                                     A;Molecule type: mRNA
A;Residues: 1-11 <GOL>
A;Cross-references: UNIPARC:UPI000017C9F3
A;Experimental source: concanavalin A-activated lymphoblast C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                          A;Reference number: PH0891; MUID:92078857; PMID:1836012
A;Accession: PH0924
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C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
C;Accession: PH0924
C;Accession: PH0924
R;Gold, D.P; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta chains in Lewis rats with experime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        clotting protein - signal crayfish
C;Species: Pacifastacus leniusculus (signal crayfish)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T-cell receptor beta chain V-D-J region (isolate 10) - rat (fragment)
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A;Molecule type: protein
A;Residues: 1-10 <KOP>
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A; Status: prelimina
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Pred. No. 2.5e+03;
1; Mismatches 1
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Pred. No. 2.8e+03;
Wismatches 0;
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Pred. No.
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                                     #text_change 10-Nov-1995
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T.; Funae,

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Biochim. Biophys. Acta 1158, 227-236, 1993
A;Title: Characterization of two P-450 isozymes placed in the rat CYP2D subfamily. A;Reference number: S39761; MUID:94072607; PMID:8251521
A;Recession: S39762
A;Molecule type: protein
A;Residues: 1-12 <OHI>A;Cross-references: UNIPARC:UPI000017C8F7
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PS0443
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R;Adelman, J.P.; Shen, K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrutta, A.; Bond R;Adelman, J.P.; Shen, K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrutta, A.; Bond Neuron 9, 209-216, 1992
A;Title: Calcium-activated potassium channels expressed from cloned complementary DNAs. A;Reference number: JH6697; MUID:92360298; PMID:1497890
A;Recession: PS0443
A;Status: nucleic acid sequence not shown
A;Atatus: nucleic acid sequence not shown
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Gross-references: UNIPARC:UPI000017BEBA
C;Comment: This potassium channel is activated by calcium.
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A;Cross-references: FlyBase:FBgn0003429
C;Keywords: alternative splicing; ion channel; potassium channel; transmembrane protein
Search completed: April
Job time : 20 secs
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|DDLV 13
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1: uniprot_sprot
2: uniprot_tremb
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Q86TJ6_HUMAN
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NUCLEOTIDE SEQUENCE.

MEDLINE=96184388; PubMed=8612793; DOI=10.1016/0014-5793 (96) 00224-4; Schmoll D., Allan B.B., Burchell A.; "Cloning and sequencing of the 5 region of the human glucose-6-phosphatase gene: transcriptional regulation by cAMP, insulin and glucocorticoids in H411R hepatoma cells.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Allan Brian B.;
Submitted (MAR-1996) to the EMBL/GenBank/DDBJ
EMBL; X96937; CAA65638.1; -; Genomic_DNA.
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NCBI_TaxID=9606;	ia; Eu	apiens	1e-6-ph	-2004	2004	,-2004	_HUMAN	E		17	17	18	18	18				18	18	18	18	18	18	
606;	Mammalia; Eutheria;	Homo sapiens (Human). Rukarvota: Metazoa: C	Glucose-6-phosphatase	(TrEMBLrel	(TrEMBLrel.	(TrEMBLrel.	Q6LAP7_HUMAN PRELIMINARY;			37.8	37.8	40.0	40.0	40.0	40.0	40.0	40.0	40.0	40.0	40.0	40.0	40.0	40.0	
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RESULT 2
Q6LCH4_RAT
ID Q6LCH4, RAT PRELIMINARY;
AC Q6LCH4,
AC Q6LCH4,
DT 05-UUL-2004 (Trembirel. 27, L4
DT 05-UUL-2004 (Trembirel. 27, L4
DT 05-UUL-2004 (Trembirel. 27, L4
         RRR OCCOORDITAC
                                        Glucose-6-phosphatase (Fragment).

Rattus norvegicus (Rat).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognau

Muridae; Murinae; Rattus.

NCBI_TaxID=10116;
NUCLEOTIDE SEQUENCE.
STRAIN=Sprague-Dawley; TISSUE=Liver;
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12 AA; 1379 MW;

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RESULT 3
O95179 HUMAN PRELIMINARY;
ID O95179;
AC O95179;
OP O1-MAY-1999 (TrEMBLrel. 10,
DT O1-MAY-1999 (TrEMBLrel. 10,
DT O1-MAY-1999 (TrEMBLrel. 19,
DT O1-DEC-2001 (TrEMBLrel. 19,
DT O1-DEC-2001 (TrEMBLrel. 19,
DT O1-DEC-2001 (TrEMBLrel. 19,
DE Glucose-6-phosphatase hydro
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordat
OC Mammalia; Eutheria; Euarcho
OC MOBI-TaxID=9606;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RX MEDLINE=99156747; PubMed=10
RA Burchell A.;
RT "Identification of a cAMP x
PHOSPHATASE hydrolytic subu
RT transcriptional regulation
RT hepatoma cells.";
RL Biochem. J. 338.457-463 (199
DR EMBL; APO5135; AAD11621.1;
FT NON TER
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SQ SEQUENCE 14 AA; 1666 FM;
       RESULT 4
Q86TJ6 HUMAN
ID Q86TJ6;
AC Q86TJ6;
DT 01-JUN-2
DT 01-JUN-2
DT 01-JUN-2
DT 101-JUN-2
DT 01-JUN-2
DE FLJ20097
OS Homo sap
OC Eukaryot
OC Mammalia
OC Homo.
OX NCBI_Tax
RN [1]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          flanking sequence.";
Diabetes 45:1563-1571(1996).
EMBL; U57552; AAB19044.1; -;
NON_TER 13 13
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Glucose-6-phosphatase hydrolytic subunit (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Identification of a cAMP response element within the glucose-phosphatase hydrolytic subunit gene promoter which is involved transcriptional regulation by cAMP and glucocorticoids in H4IIE hepatoma cells.";
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MEDLINE=99156747; PubMed=10024523;
Schmoll D., Wasner C., Hinds C.J.,
TISSUE=Peripheral Nervous System;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
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01-JUN-2003 (TrEMBLrel. 24,
01-JUN-2003 (TrEMBLrel. 24,
FLJ20097 protein.
                                                                                                                                  Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Buarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                            Q86TJ6_HUMAN
                                                                                             NCBI_TaxID=9606
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Allan B.B., Walther R.,
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RESULT 5
Q9TWT4 LUMTE
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AC Q9TWT4;
DT 01-MAY-2
DT 01-MAY-2
DT 01-JUN-2
DT 01-JUN-3
RM [1]
RA MEDLINE-RA OWNDY D.
RT "The ext
RT Determit
RL J. Bioll.
SQ SEQUENCE
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O1-MAY-2000 (TrEMBLrel. 13, Last sequence of the sequence 
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Submitted (MAR-2003) to the
EMBL; BC047757; AAH47757.1;
SEQUENCE 15 AA; 1843 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROTEIN SEQUENCE.

MEDLINE-93293879; PubMed-8514787;

MEDLINE-93293879; PubMed-8514787;

Ownby D.W., Zhu H., Schneider K., Beavis R.C., Chait B.T.,

"The extracellular hemoglobin of the earthworm, Lumbricus
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determination of subunit stoichiometry.";
J. Biol. Chem. 268:13539-13547(1993)
SEQUENCE 15 AA; 1665 MW; P9B83FE0050D
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Pred. No.
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TUF05 MOUSE P38643; 01-OCT-1994 01-OCT-1994 05-JUL-2004

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Q66874;
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EMBL; U07130; AAA74412.1; -;
NON_TER 1 1
                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                "Capsid protein gene sequence of feline calicivirus isolates LLK: further evidence for capsid protein configuration among
                                                                                                                                                                                                                                                                                                                                                                                                                                          Seal B.S., Neill J.D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUCLEOTIDE SEQUENCE. STRAIN=255;
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-I- MISCELLANEOUS: On the 2D-gel the determined pI of the protein is: 5.5, its MW is: 48 kDa.
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TISSUE=Fibroblast;
MEDLINE=95009907; PubMed=7523108;
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RESULT 9
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ID Q7M063_MOUSE PRELIMINARY;
AC Q7M063;
DT 01-MAR-2004 (TrEMBLrel. 26,
DE Calnexin (Fragment).
OE Mus musculus (Mouse).
OC Bukaryota; Metazoa; Chordat.
OC Muridae; Murinae; Mus.
OC Muridae; Murinae; Mus.
OX NCBI_TAXID=10090;
RN [1]
RN [1]
RN MEDLINE=94216347; PubMed=81
RX MEDLINE=94216347; PubMed=81
RA Lenter M., Vestweber D.;
RT Calnexin prior to integrin
RI calnexin prior to integrin
RI J. Biol. Chem. 269:12263-12
DR PIR; A53594; A53594.
FT NON_TER 15
SQ SEQUENCE 15 AA; 1693 MW;
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P14537;
01-JAN-1990 (Rel
01-JAN-1990 (Rel
01-FEB-2005 (Rel
Fibrinogen beta
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01-NOV-1996 (TrEMBLZ
01-NOV-1996 (TREMBLZ
01-NOV-1998 (TREMBLZ
Polymerase (Fragment
Feline calicivirus
Fibrinogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIINE=94216347; PubMed=8163531;
Lenter M., Vestweber D.;
The integrin chains beta-1 and alpha-6 associate with the calnexin prior to integrin assembly.;
J. Biol. Chem. 269:12263-12268(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (Mouse).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Buarchontoglires; Glires; Rodentia; Sciurogna
Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          caliciviruses.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Capsid protein gene sequence of feline calicivirus isolates LLK: further evidence for capsid protein configuration among
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seal B.S., Neill
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3; Conserv
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Last sequence update)
Last annotation update)
[Contains: Fibrinopeptide
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57.1%;
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Pred. No. 4.2e+03;
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Q47600;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              O'Neil P.B.,
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                         "A family of regulatory genes associated with modification systems.";
J. Bacteriol. 173:1367-1375(1991).
                                                                                                                                                                                                                                                                                                                                                    01-NOV-1996 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
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                                                                                                                                        Tao
                                                                                                                                                           NUCLEOTIDE SEQUENCE.
MEDLINE=91139577; PubMed=1995588;
                                                                                                                                                                                                                    NCBI_TaxID=562
                                                                                                                                                                                                                                              Enterobacteriaceae;
                                                                                                                                                                                                                                                                   Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
                                                                                                                                                                                                                                                                                     Escherichia coli.
                                                                                                                                                                                                                                                                                                                              REase protein (Fragment)
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Central domain (By similarity).

DOMAIN: A long coiled coil structure formed by 3 polypeptide chains connects the central nodule to the C-terminal domains (distal nodules). The long C-terminal ends of the alpha chains (distal nodules) are contributing a fourth strand to the coiled coil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             s Swiss-Prot entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EN European Bioinformatics Institute. There are no resti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: Fibrinogen has a polymerize into fibrin and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nonidentical chains (alpha, beta and gamma). The 2 hete are in head to head conformation with the N-termini in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      coagulation;
                                  M63619; AAA24556.1;
ER 11 11
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ch cleaves fibrinopeptides A and B from alph
thus exposes the N-terminal polymerization
the formation of the soft clot.
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                                                                                                                                      Bourne J.C.
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PS00514; FIBRIN_AG_C_DC
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hylogeny based o
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              formatics Institute. There are no restrictions content is in no way modified and this statement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1097 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (White rhinoceros) (Square-lipped rhinoceros).
Chordata; Craniata; Vertebrata; Euteleostomi;
Laurasiatheria; Perissodactyla; Rhinocerotidae;
                                                                                                                                        , Blumenthal R.M.;
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                 63175479572AB5A4 CRC64;
                                                        Genomic_DNA
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tion sites
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RESULT 13
Q9R5D6 CHRVI
ID C9R5D6_C
AC Q9R5D6;
DT 01-MAY-2
DT 01-MAY-2
DT 01-JUN-2
DE POLY(3-H
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Best Local S
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Matches 3
Q9R5D6_CHRVI PRELIMINARY;
Q9R5D6;
Q1-MAY-2000 (TrEMBLrel. 13, C
01-MAY-2000 (TrEMBLrel. 13, I
01-JUN-2000 (TrEMBLrel. 14, I
POLY(3-HYDROXYBUTYRIC acid) S
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Biochem.
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STRVN
XYLA_S
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NON_TER
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Streptomyces violaceoruber. Bacteria; Actinobacteria; A
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01-JAN-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR001998; Xylose isom.

PROSITE; PS00172; XYLOSE ISOMERASE 1; PARTIAL.

PROSITE; PS00173; XYLOSE ISOMERASE 2; PARTIAL.

Carbohydrate metabolism; Direct protein sequences to the protein sequences are the protein sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EW the European Bioinformatics Institute. There are no restr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                + +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=LMG 7183;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Magnesium; Metal-binding; Pentose ACT_SITE 5 5
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InterPro; IPR001998; Xylose_
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUBUNIT: Homotetramer.
SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: Belongs to the xylose isomerase family.
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LYTIC ACTIVITY: D-xylose = D-xylulose
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Streptomycetaceae; Streptomyces
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granule-associated 41 kDa
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0; Mismatches
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Best Local S
Matches 4
Q47556 ECOLI
Q47556;
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MEDLINE=22255706; pubMed=12368865; DOI=10.1038/nature01099;
Carlton J.M., Angiuoli S.V., Suh B.B., Kooij T.W., Pertea M.,
Silva J.C., Ermolaeva M.D., Allen J.B., Selengut J.D., Koo H.L.,
Peterson J.D., Pop M., Kosack D.S., Shumway M.F., Bidwell S.L.,
Shallom S.J., van Aken S.B., Riedmiller S.B., Feldblyum T.V.,
Cho J.K., Quackenbush J., Sedegah M., Shoaibi A., Cummings L.M.,
Florens L., Yates J.R. III, Raine J.D., Sinden R.E., Harris M.A.,
Cunningham D.A., Preiser P.R., Bergman L.W., Vaidya A.B.,
van Lin L.H., Janse C.J., Waters A.P., Smith H.O., White O.R.,
Salzberg S.L., Venter J.C., Fraser C.M., Hoffman S.L., Gardner M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q7R8X5;
01-MAR-2004
01-MAR-2004
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plasmodium yoelii yoelii.
Eukaryota; Alveolata; Api
EUKI_TaxID=73239;
                                                                                                                                                                                                                                                                                                                          preliminary data.
EMBL; AABLO1002528; E
Hypothetical protein.
SEQUENCE 9 AA; 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Genome sequence and comparative analysis parasite Plasmodium yoelii yoelii."; Nature 419:512-519(2002).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hypothetical protein. Name=PY07095;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Liebergesell M., Schmidt B., Steinbuchel A.; "Isolation and identification of granule-associated proteins relevant for poly(3-hydroxyalkanoic acid) biosynthesis in Chromatium vinosum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROTEIN SEQUENCE.
MEDLINE=93146381; PubMed=1490603; DOI=10.1016/0378-1097(92)90031-I;
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50.0%;
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66.78;
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                                                                                                                                                                                                                              Score 19;
Pred. No.
3; Mismatc
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Last sequence up
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                        PRT;
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shotgun (WGS)
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Escherichia coli Kl2.
Bacteria; Proteobacteria;
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01-NOV-1996 (TrEMBLrel.
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                                                                                                                                        SEQUENCE
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Proc. Natl. Acad. Sci.
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MEDLINE=82275057; PubMed=7051000;
Pauza C.D., Karels M.J., Navre M., Schachman H.K.;
"Genes encoding Escherichia coli aspartate transcarbamoylase:
                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=83333;
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                                                                                                                                                                                                                             MEDLINE=83195078; PubMed=6302686; ioover T.A., Roof W.D., Foltermann
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Pred. No. 2
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1: /cgm2 6/ptodata/1,
2: /cgm2 6/ptodata/1,
3: /cgm2 6/ptodata/1,
4: /cgm2 6/ptodata/1,
5: /cgm2 6/ptodata/1,
6: /cgm2_6/ptodata/1,
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/cgn2 6/ptcdata/1/iaa/H_COMB.pep:*
/cgn2 6/ptcdata/1/iaa/PCTUS_COMB.pep:*
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US-09-269-250E-20
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•	Sequence 32, Appl	Sequence 47, Appl	Sequence 14, Appl	Sequence 1, Appli	Sequence 370, App	Sequence 8, Appli	Sequence 8, Appli	Sequence 3, Appli	Sequence 72, Appl	ű	-	26,	Sequence 25, Appl	24,	Sequence 23, Appl	Sequence 22, Appl	Sequence 21, Appl

ALIGNMENTS

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FRUELL NO. 100.03.

FRUETA NO. 100.03.

APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/269,250B
CURRENT FILLING DATE: 1999-05-21

NUMBER OF SEQ ID NOS: 38
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 20
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
FEATURE:
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09489760 Patent No. 6878375
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
ORGANISM: Histocompatibility antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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Pred. No. 4.6e+05;
); Mismatches 0;
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US-09-269-250E-28
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APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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Matches 9
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PATENT NO. 6830883
GENERRAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 199-05-21
NUMBER OF SEQ ID NOS: 38
SOPTWARE: PatentIn version 3.1
SEQ ID NO 28
SPECIAL 13
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SEQ ID NO 16
LENGTH: 13
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Best Local
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GENERAL INFORMATION:
                                                                                                                                                                      ORGANISM: Unknown Organism
                                                                                                                                                                                        TYPE: PRT
                                                                                                                                       OTHER INFORMATION: Description
                                                                                                                                                       FEATURE:
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VLHDDLLEA 11
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Pred. No. 4.6e+05;
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Pred. No. 0.034;
                                                                          Score 45; DB 2;
Pred. No. 0.034;
                                                                                                                                         of Unknown Organism:HA-1+/+
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US-09-269-250E-18
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US-09-269-250E-26
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Sequence 26, Application US/09269250E
PACENT NO. 6830883
GENERAL INFORMATION:
APPLICANT: GOULMY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: GOULTMY, RISHA
APPLICANT: GOULTMY, METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT FILLNG DATE: 1999-05-21
RUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                       EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PATENTIN VEY: 2.1
SEQ ID NO 5
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                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
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Best Local
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CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
OTHER INFORMATION: Descriptio
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TYPE: PRT
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Pred. No.
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PATENT NO. 6878375

GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Roulmy Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Victor H
ITITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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US-09-269-250E-29
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US-09-269-250E-29
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US-09-489-760-14
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                                                                                                                           Sequence 29, Application US/09269250E
Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: GOLINT, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 29
       LENGTH: 9
TYPE: PRT
ORGANISM: Human
FEATURE:
NAME/KEY: MISC FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a
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Best Local Similarity
Matches 8; Conserv
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SEQ ID NO 14
LENGTH: 13
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Matches 8; Conserv
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TYPE: PRT
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ORGANISM: Unknown Organism
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Pred. No.
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Pred. No.
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Best Local Similarity
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                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
TITLE OF INVENTION: 2183-4285US
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PACENT NO. 6878375
GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
                                                                                                                                  SEQ ID NO 4
LENGTH: 9
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GENERAL INFORMATION:
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                  FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1
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NUMBER OF SEQ ID NOS: 16
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FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (2)..(7)
OTHER INFORMATION: AMINO ACIDS X REPRESENT LEUCINE OR ISOLEUCINE
OTHER INFORMATION: RESIDUES
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TYPE: PRT
ORGANISM: Histocompatibility antigen
FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (3)
                                                                                                                 TYPE: PRT
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                                                                                            ORGANISM: Histocompatibility antigen
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
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US-09-269-250E-38
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US-09-025-819-5
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Best Local Similarity bs...
Grandes 8; Conservative
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PATENT NO. 6830883
GENERAL INFORMATION:
APPLICANT: GOLIMY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PATENTIN Version 3.1
SEQ ID NO 38
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Patent No. 6225097
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                                                                                                                                       STATE: DC
COUNTRY: USA
COUNTRY: USA
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA: US/09/025,819
FILING DATE: 19-FEB-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: JP 251675
PILING DATE: 17-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: Khalilian, Houri
REGISTRATION NUMBER: 39.546
REFERENCE/DOCKET NUMBER: 10235/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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APPLICANT: Nishino, Tokuzo
APPLICANT: Koyama, Tanetoshi
APPLICANT: Sato, Yoshihiro
TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH: 8
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ADDRESSEE: KENYON &
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1500 K Street, N.W.
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Pred. No. 4.6e+05;
0; Mismatches 3
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Pred. No. 4.6e+05;
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; TOPOLOGY: linear
; MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 5:
US-09-808-126-5
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US-09-808-126-5
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Patent No. 6410280
GENERAL INFORMATION:
APPLICANT: Obata, Shusei
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/808,126
FILING DATE: 08-May-2001
CLASSIFICATION: cUnknown>
PRIOR APPLICATION NUMBER: 09/025,819
FILING DATE: CUNknown>
APPLICATION NUMBER: 09/025,819
FILING DATE: CUNKnown>
AFTORNEY/AGENT INFORMATION:
NAME: Khalilian, Houri
REGISTRATION NUMBER: 39,546
REFERENCE/DOCKET NUMBER: 10235/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 80.0%;
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                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-220-4200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE NUMBER OF SEQUENCES: 31
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STREET: 1500 K Street, N.W.
                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                           LENGTH: 7 amino acids
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Koyama, Tanetoshi
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Pred. No. 4.6e+05;
1; Mismatches 0;
                  Score 25; DB 2;
Pred. No. 4.6e+05;
 Mismatches
                                      DB 2;
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RESULT 15

US-09-803-511-5

REPUBLICATION US/03803951

PACENT NO. 6413761

REPUBLICANT: Obata, Shuse1

APPLICANT: Obata, Shuse1

REPUBLICANT: Obata, Shuse1

REPUBLICANTON: OBECAPERRYL DIPHOSPHATE SYNTHETASE GENE

MADRIES OF SEQUENCES: 12

REPUBLICANTON US/DESS: CANNOW & CENTYON

REPUBLICANTON UNMERS: PCOS/MS-DOS

CURRENDO THARE: PROPARES: 05/05/803,951

PILING DATE: 13 MG-700/803,951

PILING DATE: Obatacomb

REPUBLICANTON UNMERS: 139/803,951

PILING DATE: 139/803
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Minimum DB
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

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4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

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US-10-861-355-1
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US-11-623-176-74
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Publication No. US20040092446A1
GENERAL INFORMATION:
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
FITLE OF INVENTION: HA-1 epitopes and uses
FILE REFERENCE: 2183-6047US
CURRENT EPPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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Result No.

32 71.1 9 4 US-10-623-176-4 30 66.7 9 4 US-10-623-176-14 30 66.7 9 4 US-10-623-176-14 30 66.7 9 4 US-10-791-217-4 30 66.7 9 4 US-10-791-217-4 30 66.7 9 4 US-10-791-217-4 30 66.7 9 4 US-10-623-176-15 28.5 63.3 8 4 US-10-623-176-15 28.5 63.3 8 6 US-11-007-740-38 28.6 62.2 9 4 US-10-623-176-23 28.6 62.2 9 4 US-10-623-176-23 28.6 62.2 9 4 US-10-623-176-23 28.6 62.2 9 4 US-10-837-217-95 28.6 62.2 10 4 US-10-837-217-95 28.6 62.2 10 4 US-10-837-217-95 28.6 62.2 10 4 US-10-623-176-64 27.6 60.0 10 4 US-10-623-176-89 27.6 60.0 10 4 US-10-623-176-96	71.1 66.7 66.7 64.4 63.3 63.3 63.3 63.3 63.3 63.2 9 9 62.2 9 9 62.2 9 9 62.2 9 9 62.2 9 9 62.2 9 9 63.3 9 9 63.3 9 9 63.3 9 9 63.3 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29
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4 US-10-623-176-4 4 US-10-623-176-14 4 US-10-623-176-14 4 US-10-791-217-4 6 US-11-007-740-40 6 US-11-007-740-38 6 US-11-007-740-38 7 US-10-623-176-18 8 US-10-623-176-38 9 US-10-623-176-38 9 US-10-623-176-38 9 US-10-623-176-64 9 US-10-623-176-64 9 US-10-623-176-64 9 US-10-623-176-88	Sequence	10	10	9	13	10	9	9	ø	9	8	80	9	9	9	9	14	9
US-10-623-176-4 US-10-623-176-14 US-10-623-176-14 US-10-791-217-4 US-110-791-217-4 US-110-623-176-48 US-110-623-176-15 US-110-623-176-23 US-110-623-176-23 US-110-623-176-23 US-10-623-176-38 US-10-623-176-64 US-10-623-176-64 US-10-623-176-89 US-10-623-176-89 US-10-623-176-95	Sequence	4	4.	Ŋ	4	4.	4.	4.	4	4	Q	4.	4,	0	4	4,	4.	4
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ALIGNMENTS

thereof

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PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR PILING DATE: 1907-07-23
PRIOR PILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTIN DATE: 2000-01-24
SOFTWARE: PATENTIN VET. 2.1
SOFTWARE: PATENTIN VET. 2.1
SOFTWARE: PATENTIN VET. 2.1
                                                                                                                                  US-10-791-217-2
                                                                                                                                                                                                                                                                                                                                                                                             ; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-2
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Best Local S
Matches 9
Sequence 2, Application US/10791217
Publication No. US20040191268A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description
FEATURE:
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Pred. No. 1.7
0; Mismatches
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1.7e+06;
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US-10-861-335-1
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US-11-007-740-20
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SEQ ID NO 2
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial sequence
                        Sequence 20, Application US/11007740

Sequence 20, Application US/11007740

Publication No. US2005023350A1

GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa

TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1

FILE REFERENCE: 2799/58994-A

CURRENT APPLICATION NUMBER: US/11/007,740

CURRENT PILING DATE: 2004-12-08

PRIOR APPLICATION NUMBER: 09/269,250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/10861335 Publication No. US20050031612A1 GENERAL INFORMATION:
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Matches 9; Conserv
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Best Local Similarity
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PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/861,335
CURRENT FILLING DATE: 2004-66-04
PRIOR APPLICATION NUMBER: PCT/NLO2/00791
PRIOR FILING DATE: 2002-12-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Goulmy, Elsa A.J.M.
TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunot
TITLE OF INVENTION: of tumors
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CURRENT FILING DATE: 2004-03-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
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PRIOR WILING DATE: 1999-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: HA-1 peptide
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                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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100.0%; Pred. No. 1.7e+06;
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Sequence 76, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
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        APPLICANT: Goulmy, Bls A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments US-11-007-740-20
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US-10-623-176-43
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PRIOR FILING DATE: 2000-01-21
PRIOR PPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR PPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
RESULT 6
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SEQ ID NO 43
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Publication No. US20040092446A1
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                                                                                                                   Matches
                                                                                                                                                 Query Match
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                                                    NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                      LENGTH: 10
TYPE: PRT
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9; Conserv
                                                                                  1 VLHDDLLEA 9
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                                                                                                                 Conservative
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                                                                                                                                   100.0%; Score 45; DB 4
100.0%; Pred. No. 0.17;
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Pred. No. 1.7e+06;
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                                                                                                                       Mismatches
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PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 16
PRIOR HELLING NOS: 13
FENOTH: 13
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US-11-007-740-28
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; LOCATION: (1)..(13)
US-10-623-176-76
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Sequence 28, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
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Publication No. US20040191268A1
GENERAL INFORMATION:
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Best Local
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 76
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Best Local
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CURRENT FILING DATE: 2004-03-02
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APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
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PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: obtained from HA-1+/+ phenotype
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nes 9; Conser
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FILING DATE: 2000-01-21
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Pred. No. 0.22;
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RESULT 9

US-10-623-176-41

Sequence 41, Application US/10623176

Sequence 41, Application US/10623176

Sequence 41, Application US/10623176

Publication No. US20040092446A1

GENERAL INFORMATION:
APPLICANT: Goulny, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
TITLE OF INVENTION: HA-1 epitopes and uses thereof
UNIVERSELY CONTROLLED TO THE PROPERTY OF TH
RESULT 10
US-10-623-176-45
US-10-623-176-45
; Sequence 45, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
; APPLICANT: GOULTHY, Els A.J.M.
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CURRENT APPLICATION NUMBER: US/10/623,176

CURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: 09/489,760

PRIOR PILING DATE: 2000-01-21

PRIOR PILING DATE: 2000-01-21

PRIOR PILING DATE: 1997-07-23

PRIOR APPLICATION NUMBER: PCT/NL98/00424

PRIOR APPLICATION NUMBER: PCT/NL98/00424

PRIOR FILING DATE: 1998-07-23

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR PILING DATE: 2000-01-24
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SEQ ID NO 28
LENGTH: 13
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Best Local
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SEQ ID NO 41
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CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: SITE LOCATION: (1)..(9)
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OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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ORGANISM: Artificial Sequence
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100.0%; Pred. No. 0.22;
tive 0; Mismatches
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100.0%; Pred. No. 1.
tive 0; Mismatches
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Hunt, Donald F.

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APPLICANT: GOULMNy, EIS A.J.M.
APPLICANT: GOULMNy, EIS A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR APPLICATION NUMBER: DF 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: J998-07-23
PRIOR PILING DATE: J998-07-23
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PRIOR FILING DATE: 2000-01-21
PRIOR PILING DATE: 2000-01-21
PRIOR PILING DATE: 1997-07-23
PRIOR PELICATION NUMBER: BP 97202303.0
PRIOR PILING DATE: 1997-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: DF 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 45
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US-10-623-176-10
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US-10-623-176-45
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Matches
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LENGTH: 9
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CURRENT FILING DATE: 2003-07-18
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                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
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LOCATION: (1).
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                                                                                                                                 NAME/KEY: SITE LOCATION: (1)...
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nes 8; Conserva
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            Similarity
8; Conserv
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            Conservative
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                                      82.2%;
88.9%;
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                                      Score 37; DB 4; Length 9; Pred. No. 1.7e+06;
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RESULT 12
US-10-623-176-47
; Sequence 47, Application US/10623176
; Publication No. US20040092446A1
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; LOCATION: (1)..(9)
US-10-623-176-47
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US-10-791-217-5
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Best Local Similarity
"htches 7; Conserva
; OTHER INFORMATION: obtained from KIAA0223 partial complementary DNA US-10-791-217-5
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SEQ ID NO 47
                                                                                                                          PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5, Application US/10791217 Publication No. US20040191268A1
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APPLICANT: Hunt, Donald P.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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                                                                                                                                                                                                                                                                                                                                        APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
                                                                                                                                                                                                                                                                              FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
                                                              TYPE: PRT ORGANISM: Artificial sequence
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                                                                                                          LENGTH: 9
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Sequence 18, Application US/11007740

Publication No. US20050233350A1

GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740

CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250

PRIOR PRIOR SEQ ID NOS: 40
SOPTWARE: Patentin version 3.1
SEQ ID NO 18
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments
US-11-007-740-18
; OTHER INFORMATION: I
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (1)..(10)
US-10-623-176-44
                                                                                                        PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 1907-07-23
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EPT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: D2000-504165
PRIOR APPLICATION NUMBER: JD 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOPTWARE: Patentin Ver: 2.1
SEQ ID NO 44
LENGTH: 10
TYPE: PRT
TYPE: PRT
TYPE: PRT
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US-10-623-176-44
US-10-623-176-44
; Sequence 44, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
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Matches 8; Conserve
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US-11-007-740-18
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                              FEATURE:
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                                                                      Description of Artificial Sequence: HA-1 peptide
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Pred. No. 1.7e+06;
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Title:
Perfect score:
Sequence:
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No.
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Maximum DB seq length: 15
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Maximum Match 10
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                        Score
      Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/USOS NEW PUB.pep:*

2: /SIDSS/ptodata/2/pubpaa/USOS NEW PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/USO7 NEW PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT NEW PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/USO9 NEW PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*

7: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*

8: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*
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1 VLHDDLLEA 9
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                                                                                                                                                                                                                                                                                                                                        Length
                   100%
  US-11-010-748A-11
US-11-010-748A-12
US-11-129-143-163
US-11-129-143-165
US-11-129-143-166
US-11-129-143-162
US-11-129-143-162
US-11-129-143-162
US-11-129-143-162
US-11-004-399-115
US-10-485-788A-353
US-10-485-788A-353
US-10-485-788A-353
US-10-485-788A-353
US-10-485-788A-353
US-11-033-039-809
US-11-033-039-809
US-11-033-039-801
US-11-043-93-318
US-11-043-93-318
US-11-043-93-318
US-11-043-39-318
US-11-045-288-1
US-11-045-288-1
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Sequence 11, Appl Sequence 163, App Sequence 164, App Sequence 165, App Sequence 167, App Sequence 167, App Sequence 115, App Sequence 115, App Sequence 115, App Sequence 33, Appl Sequence 354, App Sequence 354, App Sequence 354, App Sequence 809, App Sequence 809, App Sequence 800, App Sequence 801, Appl Sequence 801, App Sequence 318, App Sequence 318, App Sequence 310, App Sequence 5, Appli Sequence 443, App Sequence 444, App Sequence 444, App
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RESULT 2 US-11-010-748A-12

Sequence 12, Application US/11010748A
Publication No. US20050244421A1
GENERRAL INFORMATION:
APPLICANT: Merck Patent GmbH
APPLICANT: STRITMATTER, Wolfgang
APPLICANT: STRITMATTER, Wolfgang
APPLICANT: SCHARM, Burkhard

STRITTMATTER, Wolfgang MOLL, Heidrun SCHARM, Burkhard

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	Query Matc Best Local Matches	NAME/F LOCATI OTHER OTHER OTHER 11-010-	ENGTI	PRIOR APP. PRIOR FIL NUMBER OF SOFTWARE:	CURRENT CURRENT PRIOR AI PRIOR FI	APPLICANT: APPLICANT: TITLE OF I	GENERAL INF APPLICANT: APPLICANT:	RESULT 1 US-11-010-748 ; Sequence 11 ; Publication		44 45	4.2 4.3	41	2 9 0	37 8	ω ω 4 τυ 4	32 33	30	N N 19 9 80 •	26 27
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o vo	100.0%; 100.0%; vative	Ć)	ens	3 6	T APPLICATION UNMBER: US/11/010,74 T PILLING DATE: 2004-12-13 APPLICATION NUMBER: PCT/EP03/06251 FILLING DATE: 2003-06-13	Heldrun I, Burkhard I: COMPOSITIONS	=			11			110		10		999		
	0			EP02013423.5 -13 3.1	US/11/010, 12-13 CT/EP03/062		ьн Wolfgang	US/11010748A 4421A1		7 US-	7 US-	-	7 US-		6 US-			1 1	us-
	Score 45; D Pred. No. 1. Mismatches	istocom		423.5	/010,748A 3/06251	AND MET	ш	48A	ALIGNMENTS	11-045- 11-045-	11-045-	11-045-	11-045-	11-045-	10-989-	11-045- 10-989-	11-045-024-	045-	11-045- 11-045-
	; DB 7; . 1.4e+05 ches 0	histocompatibility			8A	METHODS FOR			NTS	024-906 024-3923	045-024-7118 045-024-12588	024-709	024-382	US-11-045-024-61 US-11-045-024-761	US-10-989-767A-441 US-10-989-767A-547	US-11-045-024-14029 US-10-989-767A-159	US-11-045-024-5511 US-11-045-024-12585	024-607	024-5467 024-7094
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NAME/KEY: MISC FEATURE

LOCATION: (1)..(9)

OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by T

OTHER INFORMATION: ell epitopes

US-11-010-748A-12
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                                                                                                                                                                       RESULT 4
                                                                                                     Sequence 164, Application US/11129143 Publication No. US20050266518A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.1
SEQ ID NO 163
LENGTH: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 163, Application US/11129143 Publication No. US20050266518A1
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Best Local Similarity
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PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/11/010,748A CURRENT FILING DATE: 2004-12-13
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 197
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TYPE: PRT
ORGANISM: Homo sapiens
                                   APPLICANT:
                                                                                   APPLICANT: BERRY, Alan
                     APPLICANT:
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BRETZEL, Werner
HUMBELIN, Markus
LOPEZ-ULIBARRI, Rual
MAYER, Anne F.
YELISEEV, Alexei A.
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HUMBELIN, Markus
LOPEZ-ULIBARRI, Rual
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Pred. No. 1.4e+05;
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Pred. No.
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Sequence 166, Application US/11129143
Publication No. US20050266518A1
GENERAL INFORMATION:

APPLICANT: BERRY, Alan APPLICANT: BRETZEL, W.

BRETZEL, Werner HUMBELIN, Markus

SOFTWARE: PatentIn version 3.1 SEQ ID NO 166 LENGTH: 6

APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEN, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197

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RESULT 6
US-11-129-143-166
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                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1
SEQ ID NO 165
                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, We
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 57.8%;
Best Local Similarity 80.0%;
Matches 4; Conservative
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                                                                                                                                        Query Match
Best Local Similarity
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 165,
                                                                                                                                                                                                                                                                                APPLICANT: YELISERY, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                      ORGANISM: Escherichia coli
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TYPE: PRT
                                                                                                                                                                                                                      TYPE: PRT
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                                                                                            2 LHDDL 6
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HUMBELIN, Markus
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                                                                                                                                                                                                                                                                                                                                                                                            LOPEZ-ULIBARRI, Rual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/11129143
p. US20050266518A1
                                                                                                                           Conservative
                                                                                                                                      57.8%;
80.0%;
                                                                                                                        Score 26; Db /; --
Pred. No. 1.4e+05;
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                                                                                                                                                          Length 6;
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                                                                                                                             Indels
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APPLICANT: BRETZEL, Werner

APPLICANT: HUMBELIN, Markus

APPLICANT: LOPEZ-ULIBARRI, Rual

APPLICANT: LOPEZ-ULIBARRI, Rual

APPLICANT: WAYER, Anne F.

APPLICANT: YELISEV, Alexei A.

TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION

FILE REFERICE: C38435/121966

CURRENT APPLICATION NUMBER: US/11/129,143

CURRENT APPLICATION NUMBER: US/11/129,143

CURRENT FILING DATE: 2005-05-13

NUMBER OF SEQ ID NOS: 197

SOFTWARE: Patentin version 3.1

SEQ ID NO 162

LENGTH: 6

TYPE: PRT
ORGANISM: Rhizobium sp. strain NGR234

US-11-129-143-162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: HUNBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: WAYER, Anne F.
APPLICANT: YELISESV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT PILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOPTWARE: PatentIn version 3.1
SEQ ID NO 161
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-11-129-143-162
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US-11-129-143-161
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                                                         Matches
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Best Local (
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Best Local Similarity
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Publication No. US20050266518A1
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APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
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                                                                         Similarity
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                                                           Conservative
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80.0%;
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80.0%;
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80.0%;
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Pred. No. 1.4e
1; Mismatches
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Pred. No.

    Mismatches

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Pred. No.
                                                           Mismatches
                                                                         1.4e+05;
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                                                                                          Length 6;
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RESULT 11
US-10-485-788A-353
; Sequence 353, Application US/10485788A
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                                                                                                                                                                                                                                                                                                                                    Sequence 1155, Application US/11004399
Publication No. US20060053516A1
GENERAL INFORMATION:
APPLICANT: Chye, Mee Lee
APPLICANT: Li, Hong Ye
                                                                                                                                                            Query Match
Best Local Similarity
Matches 4; Conserv
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Publication No. US200660046277A1
GENERAL INFORMATION:
APPLICANT: BELYAEV, ALEXANDER SERGEEVICH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.1 SEQ ID NO 1155
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Best Local Similarity
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APPLICANT: Poon, Leo Lit Man
APPLICANTON: Genetically Modified Plants Comprising SARS-COV Viral Nucleotide
TITLE OF INVENTION: Sequences and Methods of Use Thereof For Immunization Against SA
FILE REPERENCE: 2589/73166/RDK
CURRENT APPLICATION NUMBER: US/11/004,399
CURRENT FILING DATE: 2003-12-03
PRIOR APPLICATION NUMBER: US 60/527,637
PRIOR APPLICATION DATE: 2003-12-03
NUMBER OF SEQ ID NOS: 4043
                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                        FEATURE:
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                                                                                                                2 LHDD 5
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                                                                                                                                                              Conservative
                                                                                                                                                                               53.3%;
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                                                                                                                                                                <u>,</u>
                                                                                                                                                                                 Score 24;
; Pred. No.
                                                                                                                                                                Mismatches
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98;
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RESULT 12
US-10-485-788A-354
US-10-485-788A-354
; Sequence 354, Application US/10485788A
; Publication No. US20050282743A1
; GENERAL INFORMATION:
; APPLICANT: Lu, Peter S.
; APPLICANT: Rabinowitz, Joshua D.
; APPLICANT: Schweizer, Johannes
RESULT 13
US-10-485-788A-355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
APPLICANT: Schweizer, Johannes
APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions in Cells
FILE REFERENCE: 20054-003320US
CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR FILING DATE: 2002-02-25
PRIOR FILING DATE: 2002-02-25
PRIOR FILING DATE: 2002-02-25
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
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Best Local Similarity
Workshes 3; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-485-788A-353
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                                                                                                                                             Query Match
Best Local Similarity 50.0
Workshes 3; Conservative
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SEQ ID NO 354
LENGTH: 7
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                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Molecular Interactions in Cells FILE REFERENCE: 20054-003320US CURRENT APPLICATION NUMBER: US/10/485,788A CURRENT FILING DATE: 2004-02-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Lu, Peter S.
                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                            TYPE: PRT
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                                                                                            LVHDDV 7
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Arbor Vita Corporation
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                                                                                                                                                                                                  51.1%; Score 23; DB 6; 50.0%; Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
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US-10-485-788A-355
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US-11-033-039-785
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CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
PRIOR PILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
                                                                                                                                                                        US-11-033-039-785
                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/11/033,039
CURRENT FILING DATE: 2005-01-11
PRIOR APPLICATION NUMBER: 10/245,871
PRIOR FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-07-17
PRIOR FILING DATE: 2002-07-17
PRIOR APPLICATION NUMBER: 09/396,813
PRIOR FILING DATE: 1999-09-14
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Best Local Similarity 50.0
""" Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 785, Application US/11033039

Publication No. US20060002947A1

GENERAL INFORMATION:
APPLICANT: HUMPHREYS, ROBERT

APPLICANT: XU, MINZHEN

TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES

FILE REFERENCE: REH-2017US01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1 SEQ ID NO 355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 355, Application US/10485788A Publication No. US20050282743A1
                                                                                                                                                                                                                                                          SEQ ID NO 785
                                                                                     Matches
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                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lu, Peter S.
APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions in Cells
                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 1452
                                                                                                                                                                                                                TYPE: PRT
                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                            ORGANISM: Zaire ebolavirus
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Pred. No. 1.4e+05;
3; Mismatches 0;
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RESULT 15 US-11-033-039-809 ; Sequence 809, Application US/11033039 ; Publication No. US20060002947A1

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APPLICANT: HIMPHREYS, ROBERT

APPLICANT: XI, MINZHEN

ITITLE OF INVENTION: LI-KSY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES

FILE REFERENCE: REH-2017USO1

CURRENT APPLICATION UNMERE: US/11/033,039

CURRENT PILING DATE: 2005-01-11

PRIOR APPLICATION NUMBER: 10/245,871

PRIOR PILING DATE: 2002-09-17

PRIOR FILING DATE: 2002-09-17

PRIOR PILING DATE: 2002-09-14

PRIOR FILING DATE: 2002-09-14

NUMBER OF SEQ ID NOS: 1452

SOPTWARE: Patentin version 3.3

SEQ ID NO 809

LENGTH: 99

LENGTH: 99

TYPE: PRT

ORGANISM: Zaire ebolavirus

US-11-033-039-09

Query Match
Best Local Similarity 44.4%; Pred. No. 1.4e+05;
Matches 4; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

Matches 4; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

Search completed: April 6, 2006, 17:29:30

Search completed: April 6, 2006, 17:29:30
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GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd.

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Post-processing: Minimum Match
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1: geneseqp1980
2: geneseqp2000
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7: geneseqp2003
8: geneseqp2003
9: geneseqp2003
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SUMMARIES

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m A odor 22,025mu	3	B. for	Adt73689 Human RSV	Adt72766 Human RSV		Aam47761 Ribosomal	Abp71119 E10 prote	Aag94234 Human com	Adt72767 Human RSV		Prenyl	Adt73023 Human RSV	Aaw97572 T-cell ep	Aaw99195 Minor his	Aaw97415 KIAA0223	Aaw99199 vR cell K	Adh40333 Human min	Aaw97375 HA-1 H-al	Aaw99196 Minor his	Aaw97414 KIAA0223	Aaw99198 DH cell K	Adh40334 Human min	Aaw97374 HA-1 R-al	nor hi	Description	

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25
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57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	57.1	59.5	59.5
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Abr30612	Abr30473	Abr31271	Abr30940	Abr30509	Adr73051	Abg74854	Add23767	Adg66270	Adg66271	Adg66272	Adg66269	Add23610	Aaw64577	Add23269	Abr04849	Adm18218	Adm18256	Aab81550	Abb07889	Adu38989
_	Human can	Human can	Human can	Human can	L plantar	Human CHP	Breast ca	Human pro	Human pro	Human pro	Human pro	Breast ca	Human Fac	Breast ca	Human can	Chlamydia	C trachom	Zif268 zi	Phosphoen	Mutant S1

ALIGNMENTS

RESULT 1
AAW99197
ID AAW9
XX AAW9
AC Minc
XX WO99
AN W Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease. 23-JUL-1997; 23-JUL-1998; 04-FEB-1999. Minor histocompatibility antigen HA-1 T-cell epitope #3 20-MAY-1999 AAW99197; WO9905174-A1. Homo sapiens. AAW99197 standard; peptide; 9 AA. (UYLE-) RIJKSUNIV LEIDEN (first entry) 97EP-00202303 98WO-NL000425.

Goulmy EAJM, Hunt DF, Engelhard VH;

WPI; 1999-153312/13.

A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation. HA-1 - useful to treat immune versus graft disease in bone

Disclosure; Page 15; 47pp; English.

The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency diseases

Sequence 9 AA;

DB 2;

Length 9;

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RESULT 2
AAW97374
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RESULT 3
ADH40334
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Matches 9
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                                                                                                                                           Matches
                                                                                                                                                                                                                 The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW97374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW97374 standard;
                                                                                                                                                                                                                                                                                                                                                                                     Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. detection of genetic aberrances.
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02-JUN-1998;
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  ADH40334;
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                                                                                                                                                                                                                                                                                                                                                              Claim 13; Fig 5; 59pp; English.
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                         ADH40334 standard; peptide; 9 AA.
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ilarity 100.0%;
Conservative 0
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98EP-00870125.
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Pred. No. 2e+06;
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RESULT 4
AAW99198
ID AAWS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Strittmatter W,
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Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 82; 119pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9 AA;
                                                                       DH cell
                                                                                                                                            AAW99198;
                                                                                                                                                                               AAW99198 standard; peptide; 13
                                                                                                         20-MAY-1999
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                                                                       KIAA0223 protein
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                                                                        sequence.
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AAW97414
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Best Local S
Matches 9
          Goulmy EAJM,
                                                                                                 04-FEB-1999
                                                                                                                                           Homo
                                                                                                                                                              T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune disease; neoplastic haematopoietic cell; KIAA0223 polymorphism.
                                                                           23-JUL-1998;
                                                                                                                     WO9905173-A1
                                                                                                                                                                                                                                                                               AAW97414 standard; protein; 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.
                                                                                                                                                                                                                                     20-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of severe aplastic anaemia, leukaemia, and immune deficiency diseases. The present sequence represents a KIAA0223 sequence given in the present invention
                               (UYLE-) RIJKSUNIV LEIDEN
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RESULT 6
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The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                                                                                                                                               A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
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                                                                                                                                                                                                                Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-153312/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Goulmy EAJM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunogenic peptide from minor histocompatibility antigen HA-1 - usef for inducing tolerance to transplants and prevent rejection or graft-versus-host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 38;
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                                                                                                                                                                                                       Page 32; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   aplastic anaemia; immune deficiency disease.
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                                                                                                                                                                                                                                            The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. de of genetic aberrances.
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ADH4033 ADH404 ADH404 AXX ADH404 AXX ADH404 AXX ADH40 AX
RESULT 9
AAW99199
ID AAWS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel method for providing epitopes of allelic CC variants of antigenic proteins from specific species based on single CC nucleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein flentifying, CC selecting allelic peptide/protein variants, expression product or its CC fragment encoded by DNA sequence having SNP, creating variant epitopes, CC selecting epitopes binding to MHC protein. A protein of the invention has CC cytostatic activity, and may have a use in a vaccine. The method is CC individuals from a SNP profile of one or more individuals from a CC individuals, where the SNP profile was related to disease, preferably CC cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer in the snP-related polymorphic profile. A method of the invention is useful for transplanting haematopoietic stem cells from a donor to a constitution of the progression, regression or onset of a treated disease. The present concerns used in the exemplification of the invention.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human minor histocomptibility antigen HA-1 T cell eptiope.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Strittmatter W,
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                                    20-MAY-1999
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Pred. No. 2e+0
0; Mismatches
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2e+06;
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cell KIAA0223

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RESULT 10
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Best Local :
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 23-JUL-1997;
                        23-JUL-1998;
                                                  04-FEB-1999.
                                                                          WO9905173-A1
                                                                                                 Homo sapiens
                                                                                                                           T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune dis neoplastic haematopoietic cell; KIAA0223 polymorphism.
                                                                                                                                                                           KIAA0223 polymorphism in HA-1 positive homozygous individuals.
                                                                                                                                                                                                                                                         AAW97415 standard;
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                                                                                                                                                                                                        20-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                       the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 31; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A new minor histocompatibility antigen, diseases and prevent rejection and host
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97EP-00202303
                        98WO-NL000424.
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88.9%;
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Pred. No. 1.7;
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                                                                                                                                     autoimmune disease;
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RESULT 11
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Best Local
A new minor histocompatibility antigen, {\rm HA-1} - useful to treat immune diseases and prevent rejection and host versus graft disease in bone
                                                                                                                                                                                                                                                                                                                               Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia: vaccina.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents the KIAA0223 polymorphism in HA-1 positive homozygous individuals. The specification describes an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Neoplastic haematopoietic cells presenting the peptides, in an HIA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B- and T-cells
                                                                  Goulmy EAJM,
                                                                                                                      23-JUL-1997;
                                                                                                                                                                         04-FEB-1999
                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                               23-JUL-1998;
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                                                                                                                                                                                                                                        Misc-difference
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                                                                                            (UYLE-) RIJKSUNIV LEIDEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB;
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                                                                  Hunt DF,
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                                                                                                                      97EP-00202303
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ic anaemia; immune deficiency disease.
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88.9%;
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                                                                  Engelhard
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Pred. No. 1.7;
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RESULT 12
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Best Local S
Matches 8
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medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases. Reoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B-and T-cells
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                                                                                                                                                                    Immunogenic peptide fro for inducing tolerance versus-host disease.
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                                                                               The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as
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                                                                                                                                           Claim 1; Page 39; 57pp;
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8; Conserv
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88.9%;
                                                                                                                                                                                  from minor histocompatibility antigen HA-1 - nce to transplants and prevent rejection or g \mathbf{r}
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Pred.
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                                                                            cc encoding 11 separate viral proteins: non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein L and CC eight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC eight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC expressition comprising an above peptide mixed with a pharmaceutical CC excipient or an RSV immunogenic composition comprising a recombinant CC expression vector with a nucleic acid insert encoding an above peptide. CC The present invention describes an in vitro method of detecting cytotoxic CC Tymphocytes (CTLs) that respond to a major histocompatibility complex CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (prophylactic or therapeutic) vaccine composition for a NNA-based CC immunisation, or for preparing an immune response provoking vaccine in CC the event of RSV infection (the vaccine being prepared by contacting the CC polypeptide in an immune response provoking vaccine; This cappitide sequence is a human RSV high binding affinity peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human respiratory syncytial virus; cytotoxic T lymphocyte; MHC epitope; DNA-based immunisation; RSV infection; virucio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-APR-2004; 2004WO-EP004061.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-758334/74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to novel isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 16; SEQ ID NO 919; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lasters I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ALGO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RSV
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ALGONOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Desmet J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               high affinity binding peptide SeqID 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9
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88.9%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RSV),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ytotoxic T lymphocyte; CTL;
RSV infection; virucidal; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2;
2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or purified peptides of the in particular ten RSV genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lar ten RSV genes
proteins NS-1 (also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9
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Matches 4
                      Human respiratory syncytial virus
                                                                                                                                                  Human RSV L high affinity binding peptide SeqID 1586.
                                                                          human respiratory syncytial virus; cy MHC epitope; DNA-based immunisation;
                                                                                                                                                                                                            13-JAN-2005
                                                                                                                                                                                                                                                                                                                   ADT73690 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence from preserved region VI of prenyl diphosphate synthetase (PDS), was used in the preparation of primers for the amplification of PDS DNA. Substances synthesized by PDS, i.e. polyprenyl diphosphates, are precursors of physiologically active substances, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA encoding prenyl diphosphate synthetase subunit(s) prenyl diphosphate synthetase subunit polypeptide(s),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-034975/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vitamin K and ubiquinones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prenyl diphosphate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-JUN-1996;
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IIKDDILSA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           enzymes from subunit(s).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Koike A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96JP-00154441
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57.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 27; DB 2;
Pred. No. 2e+06;
3; Mismatches
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                                                                          cytotoxic T lymphocyte; CTL;
n; RSV infection; virucidal; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shimizu N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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and methods for

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This invention relates to novel isolated or purified peptides of the CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein L and CC eight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC composition comprising an above peptide mixed with a pharmaceutical CC excipient or an RSV immunogenic composition comprising a recombinant CC expression vector with a nucleic acid insert encoding an above peptide. CC Tlymphocytes (CTLs) that respond to a major histocompatibility complex (CMHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (prophylactic or therapeutic) vaccine composition for a DNA-based CC immunisation, or for preparing an immune response provoking vaccine in the event of RSV infection (the vaccine being prepared by contacting the
                  Query Match
Best Local :
 Matches
                                                                       Sequence 9 AA;
                                                                                                                       polypeptide in an immune response-provoking amount of specific CTL). Accordingly, these peptide compositions have virucidal activity. This peptide sequence is a human RSV high binding affinity peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 16; SEQ ID NO 1586; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-758334/74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lasters
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 4.
                    Similarity
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Desmet J,
                  61.9%;
44.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stegmann T;
4
                  Score 26;
Pred. No.
   Mismatches
                  DB 8;
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hemoglobin, extracellular, component - earthworm (Lumbricus terrestris) (fragment) C;Species: Lumbricus terrestris (common earthworm) (c;Date: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997 C;Accession: S65730 R;Cushitani, K.; Higashiyama, K.; Asao, M.; Hosokawa, K. Biochim. Biophys. Acta 1292, 273-280, 1996 A;Title: Characterization of the constituent polypeptides of the extracellular hemographic in the constituent polypeptides of the extracellular hemographics. S65731; MUID:96176855; PMID:8597573 A;Accession: S65731; MUID:96176855; PMID:8597573
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C;Superfamily: cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Graves, P.E.; Kaminsky, L.S.; Halpert, J.
Biochemistry 26, 3887-3894, 1987
A;Title: Evidence for functional and structural multiplicity
A;Reference number: A26997; MUID:88000604; PMID:3651420
A;Accession: C26997
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N;Alternate names: cytochrome P450
C;Species: Rattus norvegicus (Norway rat)
C;Date: 30-Jun-1988 #sequence_revision 30-Jun-1988
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C;Species: Cucumis sativus (cucumber)
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PH0924
T-cell receptor beta chain V-D-J region (isolate 10) - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Sate: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
C;Accession: PH0924
R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
J. Exp. Med. 174, 1467-1476, 1991
J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental A;Reference number: PH0891; MUID:92078857; PMID:1836012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Matveyev, V.V.; Usmanova, A.M.; Morozova, A.V.; Collins, J.H.; Khaitlina, Biochim. Biophys. Acta 1296, 55-62, 1996
A;Title: Purification and characterization of the proteinase ECP 32 from EsA;Reference number: S71920; MUID:96350420; PMID:8765229
A;Accession: S71920
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: protein
A;Residues: 1-15 <GRA>
A;Residues: UNIPROT:Q06884; UNIPARC:UPI0000174DA7
A;Cross-references: UNIPROT:Q06884; UNIPARC:UPI0000174DA7
C;Superfamily: cytochrome P450 homology
C;Keywords: electron transfer; heme; liver; monooxygenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Graves, P.E.; Kaminsky, L.S.; Halpert, Biochemistry 26, 3887-3894, 1987 A;Title: Evidence for functional and str
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A;Experimental source: strain A2
C;Keywords: hydrolase; metalloproteinase
A;Accession: PH0924
A;Molecule type: mRNA
A;Residues: 1-11 <GOL>
A;Cross-references: UNIPARC:UPI000017C9F3
A;Experimental source: concanavalin A-activated lymphoblast C;Keywords: T-cell receptor
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A; Accession: A26997
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C;Species: Rattus norvegicus (Norway rat)
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Pred. No. 2.4e
1; Mismatches
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Pred. No. 2.4e+03;
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potassium channel protein Slo G3 - fruit fly (Drosophila melanogaster) (frag
C;Species: Drosophila melanogaster
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 16-Feb-1997
C;Accession: PS0443
R;Adelman, J.P.; Shen, K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrut
Neuron 9, 209-216, 1992
A;Title: Calcium-activated potassium channels expressed from cloned compleme
A;Reference number: JH0697; MUID:92360298; PMID:1497890
A;Accession: PS0443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Rattus norvegicus (Norway rat)
C;Date: 19-May-1994 #sequence_revision 10-Nov-1995 #text_change 10-Nov-1995
C;Accession: S39762
R;Ohishi, N.; Imaoka, S.; Suzuki, T.; Funae, Y.
Biochim. Biophys. Acta 1158, 227-236, 1993
A;Title: Characterization of two P-450 isozymes placed in the rat CYP2D sub A;Reference number: S39761; MUID:94072607; PMID:8251521
A;Accession: S39762
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S39762
A;Status: nucleic acid sequence
A;Molecule type: mRNA
A;Residues: 1-13 <ADE>
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Cell 62, 591-598, 1990
A;Title: The c4 repressors of bacteriophages P1 and P7 are
A;Reference number: 842448; MUID:90335968; PMID:1696181
A;Accession: $42449
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C;Date: 07-Sep-1994 #sequence_revision 26-May-1995 #text_change 09-Jul-2004
C;Accession: S42449
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                                                                                                                          K.Z.; Kavanaugh, M.P.; Warren, R.A.; Wu, Y.N.; Lagrutta,
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Pred. No. 3e+03;
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Pred. No.
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Pred. No. 2.7e+03;
1; Mismatches 0
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2.7e+03;
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R;Shanklin, J.; Somerville, C.
Proc. Natl. Acad. Sci. U.S.A. 88, 2510-2514, 1991
A;Title: Stearcyl-acyl-carrier-protein desaturase from higher plants
A;Reference number: A39170; MUID:91172837; PMID:2006187
A;Accession: C39170
S57584
T cell receptor
C;Species: Homo
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C39170
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N/Alternate names: light-harvesting complex LHCIIc protein
C/Species: Spinacia oleracea (spinach)
C/Date: 19-Mar-1997 #sequence_revision 23-Apr-1999 #text_change 23-Apr-1999
C/Accession: S50900
R/Walters, R.G.; Ruban, A.V.; Horton, P.
Bux. J. Biochem. 226, 1063-1069, 1994
A/Title: Higher plant light-harvesting complexes LHCIIa and LHCIIc are bound A/Reference number: S50900; MUID:95112835; PMID:7813461
A/Accession: S50900
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S50900
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C;Keywords: oxidoreductase
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                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Persea americana (avocado)
C;Date: 24-Jan-1992 #sequence_revision 24-Jan-1992
C;Accession: C39170
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C;Superfamily: chlorophyll a/b-binding protein
C;Keywords: chlorophyll; chloroplast; light-harvesting complex;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPARC:UPI000017BEBA
C;Comment: This potassium channel is activated by calcium.
C;Genetics:
C;Genetics:
A;Gene: FlyBase:slo
A;Cross-references: FlyBase:FBgn0003429
C;Keywords: alternative splicing; ion channel; potassium channel; transmembrane
                                                                                                                                                                                                                                                                                                                         A; Status: preliminary
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C;Date: 19-Oct-1995 #sequence_revision 17-Nov-1995 #text_change 05-Nov-1999
C;Accession: S57584
R;Burrows, S.R.; Silins, S.L.; Moss, D.J.; Khanna, R.; Misko, I.S.; Argaet, V.P.
submitted to the EMBL Data Library, June 1995
A;Description: T cell receptor repertoire for a viral epitope in humans is diversified A;Reference number: S57494
A;Accession: S57584
A;Status: preliminary
A;Atatus: preliminary
A;Molecule type: mRNA
A;Residues: 1-15_4BURA
A;Cross-references: UNIPARC:UPI0000116747; EMBL:Z49956; NID:g887466; PIDN:CAA90227.1; PC;Keywords: T-cell receptor
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S68004
                                                                                                                                                                                                                                                                                       FEBS Lett. 375, 159-161, 1995
A;Title: Hucolin, a new corticosteroid-binding protein from human plasma with structural A;Reference number: S68004; MUID:96087107; PMID:7498469
A;Accession: S68004
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-7 <EBG>
A;Cross-references: UNIPARC:UPI000017C164
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C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
C;Accession: S68004
R;Edgar, P.P.
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Search completed: April Job time : 20 secs
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Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 3; Conservative 0; Mismatches 0;
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1: uniprot_sprot:*
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Q7M62 ECOLI
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Q9UC60 HUMAN
Q9T27 HUMAN
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Muroldea; Muridae; Murinae; Mus.
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"Newcastle disease virus fusion and motiff as markers for viral lineage Avian Pathol. 32:361-373(2003).
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AC Q7M063;
AC Q7M063;
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DE Calnexin (Fragment).
OS Mus musculus (Mouse).
OC Eukaryota, Metazoa; Chordat
OC Muridae; Murinae; Mus.
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Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELLANEOUS: On the 2D-gel the dete
protein is: 6.4, its MW is: 48.4 kDa.
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RESULT 8
Q7M062 MOUSE
Q7M062 MOUSE PRELIMINARY;
AC Q7M062;
DT 01-MAR-2004 (TYEMBLrel. 26,
DT 01-MAR-2004 (TYEMBLrel. 26,
DT 01-MAR-2004 (TYEMBLrel. 26,
DE COLlagen type I (Fragment).
OS Mus musculus (Mouse).
OC Mukaryota; Metazoa; Chordata
OC Muridae; Murinae; Mus.
OX MUSI_TAXID=10090;
RN [1]
RP PROTEIN SEQUENCE.
RX MIZUNO M., Kitafima T., Tom:
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P82560
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Best Local
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01-MAR-2004 (TrEMBLrel 26,
01-MAR-2004 (TrEMBLrel 26,
Collagen type I (Fragment).
                                                                                                                                     Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Euarchonto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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Submitted (MAY-2000) to Swiss-Prot.
-!- MASS SPECTROMETRY: MW=30142.85; METHOD=Electrospray.
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STRAIN=JRS4;
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01-OCT-2003 (TrEMBLrel. 25, I
Unknown protein from 2D-page
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P82560;
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NON_TER
SEQUENCE
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J. Biol. Chem. 269:12263-12268(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NON TER
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MEDLINE=94216347; PubMed=8163531;
Lenter M., Vestweber D.;
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VanBogelen R.A.;
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3; Conser
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15 AA;
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    PubMed=9244181; DOI=10.1016/0167-4889(95)00151-4; ma T., Tomita M., Kuboki Y.;
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15
1693 MW;
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Euarchontoglires; Glires; Rodentia; Sciurogna
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37.5%;
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-page (Fragment).
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Pred.
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6.5e+03;
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RESULT 9
QSUMM5 SULSO
AC QSUMM5;
DT 01-MAY-2000 (
DT 01-MAY-2000 (
DT 01-JUN-202 (
DE Ribosomal pro
OS Sulfolobus so
OC Archaea; Cren
OC Archaea; Cren
OC Archaea; Cren
CR INFOIDINE SEQUE
RP PROTEIN SEQUE
RX MEDLINE 92042
RA Casiano C., T
RT "Protein topc
RT linking with
RT neighbors.";
RL J Biol. Chem
FT NON_TER
SQUENCE 1(
RESULT 10
QUUNLS HUMAN
ID QUUNLS H
AC QUUNLS;
DT 01-MAY-2
DT 01-MAY-2
DT 01-MAY-2
DT 01-MAY-2
DT 01-MAY-2
OS HOMO SAP
OC ENKARYOL
OC MAMMAIA
OC HOMO.
OX NCBI_TAX
RN [1]
RP NUCLEOTI
RA Carr 1.M
RA Meredith
RL Submitte
DR EMBL; AFF
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SQUENCE
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Q9UNLE,
Q9UNLE,
01-MAY-2000 (TrEMBLrel. 13, C
01-MAY-2000 (TrEMBLrel. 13, L
01-MAY-2000 (TrEMBLrel. 13, L
APC2 protein (Fragment).
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neighbors.";
J. Biol. Chem. 2f
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NON TER
NON TER
   SEQUENCE
                          Submitted (DEC-1998) to the EMBL; AF110338; AAD29275.1; NON_TER 1
                                                             NUCLEOTIDE SEQUENCE.
Carr I.M., Markham A
Meredith D.M.;
                                                                                                                                  Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                   Name=APC2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Casiano C., Traut R.R.;
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01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "The osteoblastic MC3T3-E1 cells synthesized C-terminal propeptide type I collagen, which promoted cell-attachment of osteoblasts."; Biochim. Biophys. Acta 1310:97-102(1996).
PIR: S62675; S62675.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sulfolobus solfataricus
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15 AA; 1729 MW;
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                                                                                                                                Chordata; Craniata
Euarchontoglires;
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-; Genomic_DNA.
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Pred. No. 9.5e
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ID Q4766
ID Q4766
DT Q1-NC
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Q47602;
Q47602;
O1-NOV-1996 (TrEMBLrel. 01,
O1-NOV-1996 (TrEMBLrel. 01,
O1-DEC-2001 (TrEMBLrel. 19,
REase protein (Fragment).
Name=REase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAY-1999
01-MAY-1999
01-MAY-1999
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J. Bacteriol. 173:1367-1375(1991).
EMBL; M63620; AAA24558.1; -; Genomic_DNA.
NON_TER 11 11
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MEDLINE=91139577; PubMed=1995588;
Tao T., Bourne J.C., Blumenthal R.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Proteobacteria; Gammaproteobacteria;
Enterobacteriaceae; Escherichia.
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                                                                                                                                                                                                                                                                                                                streptococci.";
Mol. Microbiol. 31:0-0(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Streptococcus.
NCBI_TaxID=1314;
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Bacteria; Firmicutes; I
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Gukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicotyledons;

rosids; eurosids I; Cucurbitales; Cucurbitaceae; Cucumis.
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Name=PSAM;
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05-JUL-2004
                                         PROTEIN SEQUENCE.

MEDILINE=95392215; PubMed=7663175; DOI=10.1006/prep.1995.1049;
De Pauw P., Neyt C., Vanderwinkel E., Wattiez R., Falmagne P.;
Per Pauw P., Neyt C., Vanderwinkel E., Wattiez R., Falmagne P.;
Pharacterization of human serum N-acetylmuramyl-L-alanine ami
purified by affinity chromatography.";
Protein Expr. Purif. 6:371-378(1995).
GO; GO:0008745; P:N-acetylmuramyl-L-alanine amidase activity.
GO; GO:0001519; P:peptide amidation; NAS.
GO; GO:0001519; P:peptide amidation; NAS.
                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2000 (TrEMBLrel. 24, Last annotation update)
N-acetylmuramyl-L-alanine amidase (EC 3.5.1.28) (Fragment).
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Chloroplast; Direct protein
Photosystem I; Thylakoid.
NON_TER 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This Swiss-Prot entry is copyright. It is produced through a collaboratic between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Iwasaki Y., Ishikawa H., Hibino T., Takabe T.;
"Characterization of genes that encode subunits of cucumber PS I complex by N-terminal sequencing.";
Biochim. Biophys. Acta 1059:141-148(1991).
-!- FUNCTION: Essential for the activity of NADP photoreduction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q9UC60 HUMAN PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=3659;
                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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Ol-MAY-2000 (TrEMBLrel. 14, Last amoutation update)
DT
Ol-MAY-2000 (TrEMBLrel. 14, Last amoutation update)
DT
Ol-JUN-2000 (TrEMBLrel. 14, Last amoutation update)
DE
BCL2-interactive cell death susceptibility regulator (Fragment).
OS
Homo sapiens (Human).
OC
ENCATYOTIS; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC
Mammalia; Butheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
OC
Homo.
OX
NCBI_TaxID=9606;
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[1]
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McIjerink J.P., Smetsers T.F., Sloetjes A.W., Linders E.H.,
RA
Meijerink J.P., Smetsers T.F., Sloetjes A.W., Linders E.H.,
RA
Mensink E.J.;
RT
malignancies, ";
RL
Leukemia 9:1028-1832(1995).
SQ
SEQUENCE 15 AA; 1860 MW; 4F9238D6FE9743DB CRC64;
Search completed: April 6, 2006, 17:02:57 Job time : 116.333 secs
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MEDLINE=96071131; PubMed=7475270;
Medijerink J.P., Smetsers T.F., Sloetjes A.W., Linders E.H.,
Medijerink J.P., Smetsers T.F., Sloetjes A.W., Linders E.H.,
Mensink E.J.;
"Bax mutations in cell lines derived from hematological
malignancies.";
Leukemia 9:1826-1832(1995).
Leukemia 9:1826-1832(1995).
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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US-09-489-760-16
US-09-489-760-2
US-09-489-760-3
US-09-269-250E-29
US-09-489-760-3
US-09-217-609A-11
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20,
                                                                                                                                                                              Sequence 5, Application US/09489760
Patent No. 6678375
GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLS OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 198-07-23
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                             NUMBER OF SEQ ID NOS: 16
SOPTWARE: PatentIn Ver. 2.1
SEQ ID NO 5
LENGTH: 9
TYPB: PRT
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ORGANISM: KIAA0223 partial complementary DNA
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ALIGNMENTS

; OTHER INFORMATION: Description of Artificial Sequence: Exon US-09-269-250E-18 APPLICATION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1 TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1 FILE REFERENCE: 58994 CURRENT APPLICATION NUMBER: US/09/269,250E CURRENT FILING DATE: 1999-05-21 NUMBER OF SEQ ID NOS: 38 SOFTWARE: Patentin version 3.1 SEQ ID NO 18 Query Match Best Local GENERAL INFORMATION: Sequence 18, Application US/09269250E Patent No. 6830883 LENGTH: 9 TYPE: PRT ORGANISM: Artificial Sequence 1 VLRDDLLEA 9 |||||||| 1 VLRDDLLEA 9 h 100.0%; Score 42; DB 2; Similarity 100.0%; Pred. No. 4.6e+05; 9; Conservative 0; Mismatches 0; Length 9; Indels Fragments 0; Gaps 0

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GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
ITILE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILLING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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US-09-269-250E-26
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APPLICANT: GOULMY, Elsa
APPLICANT: GOULMY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 5894
CURRENT FILING DATE: 199-05-21
RUMBER OF SEQ ID NOS: 38
NUMBER OF SEQ ID NOS: 38
                                                                                                                       ; OTHER INFORMATION: Description of Unknown Organism:HA-1-/-US-09-489-760-14
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SEQ ID NO 26
LENGTH: 13
TYPE: PRT
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SEQ ID NO 14
LENGTH: 13
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ORGANISM: Unknown Organism
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100.0%; Pred. No. 4.6e+05;
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Pred. No.
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Patent No. 6878375

Patent No. 6878375

GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: GOLINY, Elsa A.J.M

APPLICANT: Hunt, Donald F

APPLICANT: Hard, Victor H

APPLICANT: Hard, Victor H

TITLE OF INVENTION: The HA-1 Antigen

TITLE OF INVENTION INDEER: US/09/489,760

CURRENT FILING DATE: 2000-01-21

CURRENT FILING DATE: 1998-07-23

NUMBER OF SEQ ID NOS: 16

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US-09-269-250E-20
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 20
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GENERAL INFORMATION:
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Matches 8
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GENERAL INFORMATION:
APPLICANT: GOULMY, Elsa
APPLICANT: GOULMY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
TITLE REFERENCE: 58994
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                            Sequence 28, Application US/09269250E Patent No. 6830883
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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILLNG DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 16
LENGTH: 13
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                                                                                                   APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOPTWARE: Patentin version 3.1
SEQ ID NO 29
LENGTH: 9
TYPE: NEW TYPE APPLICATION NUMBER OF SEQ ID NO 29
LENGTH: 9
                                                                                                                                                                                                                                                                 Sequence 29, Application US/09269250E Patent No. 6830883
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                    ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Unknown Organism FEATURE:
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Pred. No. 0.
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Pred. No. 0.29;
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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
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US-09-217-609A-11
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US-09-489-760-1
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Best Local Similarity
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APPLICANT: MURAMA
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APPLICANT:
APPLICANT:
            COUNTRY: US
ZIP: 20036
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
SOFTWARE: WordPerfect 6.1 for Windows
                                                                                                                                                                                            APPLICANT: KOYAMA, Tanetoshi
APPLICANT: SHIMIZU, Naoto
APPLICANT: CHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 9
CURRENT APPLICATION DATA:
                                                                                                                                   STATE:
                                                                                                                                                 CITY: Washington
                                                                                                                                                                STREET:
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1025 Connecticut Avenue,
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OGURA, Kyozo
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88.9%;
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05
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US-08-873-235B-11
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Patent No. 6174715
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                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
SOFTWARE: WordPerfect 6.1 for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/873,235B
FILING DATE: 11-Jun-1997
                                                                                                          APPLICATION NUMBER: JP 154441/1996
FILING DATE: 14-Jun-1996
ATTORNEY/AGENT INFORMATION:
NAME: TOFFENETI, Judith L.
REGISTRATION NUMBER: 39,048
REFERENCE/DOCKET NUMBER: 10235/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-429-1776
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APPLICANT: MURAMA
APPLICANT: KOIKE,
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INFORMATION FOR SEQ ID NO:
                                                 TELEFAX: 202-429-0796
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: 10235/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-429-1776
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ATTORNEY/AGENT INFORMATION:
NAME: TOFFENETTI, Judith L.
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: OGURA, Kyozo
APPLICANT: KOYAWA, Tanetoshi
APPLICANT: SHIMIZU, Naoto
APPLICANT: CHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
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PRIOR APPLICATION DATA:
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LENGIM: y ....
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les 4; Conserv
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TOPOLOGY: linear
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1025 Connecticut Avenue, NW -
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ilarity 57.1%;
Conservative
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Pred. No. 4.6e+05;
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Best Local Similarity
Watches 5; Conserva
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US-09-269-250E-38
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                                                                                                 US-09-972-115A-64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/129,123
PRIOR FILING DATE: 1999-04-13
NUMBER OF SEQ ID NOS: 64
SOPTWARE: PATENTIN VERSION 3.1
SEQ ID NO 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn version 3.1
SEQ ID NO 38
LENGTH: 8
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                                   Matches
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CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
CURRENT FILING DATE: 1999-05-21
KUMBER OF SEQ ID NOS: 38
                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/972,115A
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: US 60/128,577
PRIOR FILING DATE: 2000-04-10
                                                                                                                                                                                                                                                                                                                                            APPLICANT: Geron Corporation
APPLICANT: Gregg, Morin B.
APPLICANT: Walter, Funk D.
APPLICANT: Mater, Funk D.
                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: A Second Mammalian Telomerase FILE REFERENCE: 080/003C
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                                                                                                               LENGTH: 11
TYPE: PRT
ORGANISM: Homo sapiens
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   3 RDDLLE 8
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                                                    Similarity
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57.1%;
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100.0%; Pred. No. 4.6e+05;
tive 0; Mismatches 0;
                                                  Score 23; DB 2;
Pred. No. 1.7e+02;
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Pred. No. 4.6e+05;
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US-08-962-284-8

Sequence 8, Apsilication US/08962284

Patent No. 5985608

Patent No. 5985608

GENERAL INFORMATION:

APPLICANT: Land, Bliadbeth J.

APPLICANT: Pestconjamasp, Kersi N.

APPLICANT: Mulfkulle, Julia D.

TILE OF INVERTION: AUD NUCLEIC ACIDS ENCODING THE SAME

CORRESSEES: PLANCESSEES:

ADDESSEES: PLANCESSEES:

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Minimum DB
Maximum DB
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No.
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Maximum Match 100%
Listing first 45 summaries
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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US-10-623-176-10
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US-11-077-740-14
US-10-623-176-42
US-11-007-740-26
US-10-623-176-42
US-10-623-176-42
US-10-623-176-2
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US-10-623-176-43
US-10-623-176-74
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US-10-623-176-74
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US-10-623-176-43
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                    Sequence
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RESULT 2 US-10-791-217-5

Sequence 5, Application US/10791217
Publication No. US20040191268A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H

Query M Best Lo Matches Qy	; SEQ ID NO ; EEQ ID NO ; LENGTH: TYPE: PR ; ORGANISM ; PEATURE: ; PEATURE: ; PEATURE: ; NAME/KEY ; LOCATION US-10-623-17	CURRENT A CCURRENT F PRIOR APP	RESULT 1 US-10-623-176-10 Sequence 10, A Publication No GENERAL INFORM APPLICANT: G APPLICANT: B APPLICANT: B APPLICANT: B APPLICANT: B APPLICANT: B	ፋፋፋፋ ሪይፋን	22266666666666666666666666666666666666
atch cal Simi 9; 1 VLR 1 VLR	6. 4 . 4 . 4	REFERENCE: 2183-60470S INT APPLICATION NUMBER: INT FILING DATE: 2003-01-01-01-01-01-01-01-01-01-01-01-01-01-	SULT 1 -10-623-176-10 Sequence 10, Application No. US20 GENERAL INFORMATION: APPLICANT: Goullmy, APPLICANT: Hunt, DO APPLICANT: Engelha TITLE OF INVENTION:		N N N N N N N N N N N N N N N N N N N
Similarity 9; Conservat VLRDDLLEA 9	0. <	TEPEKENCE: 2183-6047US TEPEKENCE: 2183-6047US TEPEKENCE: 2003-07-18 APPLICATION NUMBER: 09/489, FILING DATE: 2000-01-21 APPLICATION NUMBER: EP 9720 FILING DATE: 1997-07-23 APPLICATION NUMBER: PCT/NL9 FILING DATE: 1998-07-23 APPLICATION NUMBER: JP 2000 FILING DATE: 1908-07-23 APPLICATION NUMBER: JP 2000 FILING DATE: 2000-01-24 OF SEQ ID NOS: 77		4444	U U U U U U U U U U U U U U U U U U U
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0%; Score 42; DB 4; Length 0%; Pred. No. 1.7e+06; 0; Mismatches 0; Inde	ce ion of Artificial Sequence:	EFEKENCE: 2183-6047US T APPLICATION NUMBER: US/10/623,176 T FILING DATE: 2003-07-18 APPLICATION NUMBER: 09/489,760 FILING DATE: 2000-01-21 APPLICATION NUMBER: EP 97202303.0 FILING DATE: 1997-07-23 APPLICATION NUMBER: PCT/NL98/00424 FILING DATE: 1998-07-23 APPLICATION NUMBER: PCT/NL98/00424 FILING DATE: 1998-07-23 APPLICATION NUMBER: JP 2000-504165 FILING DATE: 2000-01-24 OP SEQ ID NOS: 77	tion US/10623176 040092446A1 Els A.J.M. onald F. rd, Victor H. HA-1 epitopes and uses thereof	US-10-271-708-16 US-10-623-176-4 US-10-503-135-26 US-10-503-135-64 ALIGNMENTS	US-10-623-176-64 US-10-623-176-5 US-10-623-176-47 US-10-623-176-62 US-10-623-176-60 US-10-623-176-6 US-10-623-176-8 US-10-623-176-8 US-10-623-176-8 US-10-623-176-8 US-10-623-176-39 US-10-153-344-11 US-09-834-765-349 US-10-623-176-39 US-10-623-176-39 US-10-623-176-39 US-10-623-176-39 US-10-623-176-39 US-10-805-650-90
ngth 9; Indels 0; Gaps	: HA-1 peptide			26, 64,	Sequence 64, P Sequence 5, Ap Sequence 62, P Sequence 60, P Sequence 60, Ap Sequence 64, Ap Sequence 84, Ap Sequence 11, P Sequence 319, P Sequence 319, P Sequence 90, P
0;	·			Appl Appli Appli Appl Appl	Appl Appl Appl Appl Appl Appl Appl Appl

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Sequence 44, Application US/10623176

Publication No. US20040092446A1

GENERAL INFORMATION:

APPLICANT: GOLIMY, Els A.J.M.

APPLICANT: Hunt, Donald F.

APPLICANT: Engelhard, Victor H.

FILE REFERENCE: 2183-6047US

CURRENT APPLICATION NUMBER: US/10/623,176

COURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: 09/489,760

PRIOR APPLICATION NUMBER: EP 97202303.0
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; TYPE: PRT
; ORGANISM: Artificial sequence
; FEATURE:
; FEATURE:
; OTHER INFORMATION: obtained from KIAA0223 partial complementary
US-10-791-217-5
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                                                                                                                                                                                                                                                                            RESULT 4
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Best Local S
Matches 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.1 SEQ ID NO 18
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CURRENT FILLING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: WETHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: The HA-1 Antigen FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial Sequence
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Pred. No. 1.7
0; Mismatches
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GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR APPLICATION NUMBER: EP 97202303.0
                                                                                                                                ; LOCATION: (1)..(12) US-10-623-176-65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-623-176-65; Sequence 65, Application US/10623176; Publication No. US20040092446A1
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                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 65
LENGTH: 12
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SEQ ID NO 44
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                                                                     Matches
                                                                                                  Query Match
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PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
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NAME/KEY: SITE
LOCATION: (1)..
                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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NUMBER OF SEQ ID NOS: 77
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TYPE: PRT
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                        NAME/KEY: SITE
                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                TYPE: PRT
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es 9; Conserv
                                                                                    Local Similarity
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                  1 VLRDDLLEA 9
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VLRDDLLEA
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Pred. No. 0.26;
                                                                   Score 42; DB 4
Pred. No. 0.31;
Mismatches
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                                                                                                        DB 4;
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US-10-623-176-74 ; Sequence 74, Application US/10623176 ; Publication No. US20040092446A1

RESULT 6

GENERAL INFORMATION:

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FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 14
LENGTH: 13
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                              S
                                                                                                                                                           ; TYPE: PRT
; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: obtained from HA-1-/- phenotype
US-10-791-217-14
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US-10-623-176-74
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Best Local Similarity
Weatches 9; Conserve
                                                                           Query Match
Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
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CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
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APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PCT/NL98/00424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: KIAA0223 OTHER INFORMATION: sequence derived from a presumed HA-1 negative OTHER INFORMATION: individual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 13
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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                              VLRDDLLEA 9
VLRDDLLEA 11
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No. US20040191268A1
                                                                         100.0%; Score 42; DB 4; Length 13; ilarity 100.0%; Pred. No. 0.34; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 0.34;
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APPLICANT: GOLLLW, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT EILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILLING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILLING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: J000-504165
PRIOR FILLING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: J000-504165
PRIOR FILLING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
NUMBER OF SEQ ID NOS: 77
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Sequence 26, Application US/11007740

Sequence 26, Application US/11007740

Publication No. US20050233350A1

GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa
FIITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1

FILE REFERENCE: 2799/58994-A

CURRENT APPLICATION NUMBER: US/11/007,740

CURRENT APPLICATION NUMBER: 09/269,250

PRIOR APPLICATION NUMBER: 09/269,250

PRIOR APPLICATION NUMBER: 09/269,250

PRIOR FILING DATE: 1999-05-21

NUMBER OF SEQ ID NOS: 40

SOFTWARE: Patentin version 3.1

SEQ ID NO 26
                                                                                                                                                    ; FEATURE:
; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-42
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Best Local Similarity
Thehes 9; Conserve
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                                                        Best Loc
Matches
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SEQ ID NO 42
LENGTH: 9
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                                                                                                        Query Match
                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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                                                                               Local Similarity
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VLRDDLLE 8
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                                                           Conservative
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                                                      90.5%; Score 38; DB 4; L
100.0%; Pred. No. 1.7e+06;
live 0; Mismatches 0;
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US-10-623-176-46

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                                                                                                                              PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
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Best Local
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-10-623-176-46
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT PEPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: ECT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
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PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: HA-1 epitopes and uses thereof FILE REFERENCE: 2183-6047US CURRENT APPLICATION NUMBER: US/10/623,176 CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
                                                          TYPE: PRT
CRGANISM: Artificial Sequence
OTHEK INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE:
                                              FEATURE:
                                                                                                                LENGTH: 9
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100.0%; Pred. No. 1.7e+06;
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GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa A.J.M

APPLICANT: Goulmy, Elsa A.J.M

APPLICANT: Goulmy, Clark

APPLICANT: Hunt, Donald F

APPLICANT: Engelhard, Victor H

TITLE OF INVENTION: The HA-1 Antigen

FILE REFERENCE: 2183-4285US

CURRENT APPLICATION NUMBER: US/10/791,217

CURRENT FILING DATE: 2004-03-02

PRIOR APPLICATION NUMBER: US/09/489,760

PRIOR PILING DATE: 2000-01-21

PRIOR FILING DATE: 1998-07-23

NUMBER OF SEQ ID NOS: 17

COPTMINE: DATE: 1998-07-23
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US-10-791-217-2
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; LOCATION: (1)..
US-10-623-176-2
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; OTHER INFORMATION: HA-1 peptide US-10-861-335-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-861-335-1
                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: GOULNY, Blea A.J.M.
APPLICANT: GOULNY, Minor histocompatibility antigen HA-1: target antigen for immunor TITLE OF INVENTION: of tumors
FILE REFERENCE: 2183-6479US
CURRENT APPLICATION NUMBER: US/10/861,335
CURRENT FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: PCT/NL02/00791
PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR APPLICATION NUMBER: EP 01204704.9
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SEQ ID NO 2
LENGTH: 9
                                                                                                                       PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
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Publication No. US20040191268A1
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                                       TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                     LENGTH:
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Local Similarity 88.9%;
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Pred. No. 1.7e+06;
0; Mismatches 1,
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APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR FILING DATE: 299-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin version 3.1
SEQ ID NO 20
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:

                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 09/489,760
PRIOR PILLING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILLING DATE: 1997-07-23
PRIOR FILLING DATE: 1998-07-23
PRIOR FILLING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILLING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILLING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 43
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US-10-623-176-43
US-10-623-176-43
j Sequence 43, Application US/10623176
j Publication No. US20040092446A1
j GENERAL INFORMATION:
; OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide ; FEATURE: ; NAME/KEY: SITE ; LOCATION: (1)..(10) US-10-623-176-43
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US-11-007-740-20
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Matches 8; Conserv
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Best Local Similarity
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                      LENGTH: 10
TYPE: PRT
ORGANIEM: Artificial Sequence
FEATURE:
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Pred. No. 1.7e+06;
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Result
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Maximum Match 100%
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1: /SIDSS/ptodata/2/pubpaa/US08 NEW PUB.pep:*

2: /SIDSS/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/US03_NEW_PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

7: /SIDSS/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

8: /SIDSS/ptodata/2/pubpaa/US0_NEW_PUB.pep:*
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length: 15
Match
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Copyright (c) 1993 - 2006 Biocceleration
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US-11-019-027-17
US-11-096-706-52
US-11-225-686-211
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US-11-225-686-211
US-11-202-009-211
US-11-202-009-211
US-11-202-009-211
US-11-202-009-216
US-10-989-767A-219
US-10-989-767A-259
US-10-989-767A-268
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US-11-129-143-179
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Sequence 11,
Sequence 17,
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565,
159,
          RESULT 2
US-11-010-748A-11
; Sequence 11, Application US/11010748A
; Publication No. US20050244421A1
; GENERAL INFORMATION:
; APPLICANT: Merck Patent GmbH
; APPLICANT: STRITTMATTER, Wolfgang
; APPLICANT: MOLL, Heidrun
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ALIGNMENTS	US-11-225-686-3877	US-11-225-686-2954	US-11-172-740-1404	US-11-126-841A-8	US-11-106-932-84	US-10-522-297-13	US-10-522-297-12	US-10-718-264-162	US-10-718-264-162	US-10-353-783-71	US-11-127-677-130	US-11-004-399-1909	US-11-254-419-67	US-11-254-419-60	US-10-927-634-63	US-10-927-435-63	US-11-021-305-66	US-11-021-305-65	US-11-021-305-62	US-11-096-706-164
	Sequence 3877, Ap	Sequence 2954, Ap	Sequence 1404, Ap	Sequence 8, Appli	Sequence 84, Appl	Sequence 13, Appl	Sequence 12, Appl	Sequence 162, App		Sequence 71, Appl	Sequence 130, App	8	•	Sequence 60, Appl	•	Sequence 63, Appl	Sequence 66, Appl	Sequence 65, Appl	Sequence 62, Appl	Sequence 164, App

RESULT 1 US-11-010-748A-12 Query Match Best Local S Matches 9 Sequence 12, Application US/11010748A Publication No. US20050244421A1 GENERAL INFORMATION: APPLICANT: Merck Patent GmbH APPLICANT: STRITTMATTER, Wolfgang SOFTWARE: PatentIn version 3.1 SEQ ID NO 12 TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR MODULATING IMMUNE RESPONSE FILE REPERENCE: MER-136 CURRENT APPLICATION NUMBER: US/11/010,748A CURRENT FILING DATE: 2004-12-13 PRIOR APPLICATION NUMBER: PCT/EP03/06251 PRIOR FILING DATE: 2003-06-13 PRIOR PPLICATION NUMBER: EP02013423.5 PRIOR APPLICATION NUMBER: EP02013423.5 PRIOR APPLICATION SUMMER: EP02013423.5 PRIOR APPLICATION STEE EP02013423.5 PRIOR APPLICATION SUMMER: EP02013423.5 PRIOR PILING DATE: 2002-06-13 NUMBER: OF SEQ ID NOS: 926 APPLICANT: MOLL, Heidrun APPLICANT: SCHARM, Burkhard ORGANISM: Homo sapiens FEATURE: NAME/KEY: NAME/KEY: LOCATION: (1)..(9) OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by TOTHER INFORMATION: ell epitopes TYPE: PRT ENGTH: 1 VLRDDLLEA 1 VLRDDLLEA Similarity 9; Conserv 100.0%; Score 42; DB 7; ilarity 100.0%; Pred. No. 1.4e+05; Conservative 0; Mismatches 0; 9 9 DB 7; Length 9; Indels 0; Gaps 0

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SCHARM, Burkhard

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US-11-06-706-52

; Sequence 52, Application US/11096706

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PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
SOPTWARE: Patentin version 3.1
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CURRENT APPLICATION NUMBER: US/11/010,748A
CURRENT FILING DATE: 2004-12-13
                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/605,902
PRIOR FILING DATE: 2004-08-31
PRIOR APPLICATION NUMBER: 60/531,714
PRIOR FILING DATE: 2003-12-22
NUMBER OF SEQ ID NOS: 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: A-890A
CURRENT APPLICATION NUMBER: US/11/019,027
CURRENT FILING DATE: 2004-12-21
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APPLICANT: CARTER, Paul
APPLICANT: MARTIN, Francis H.
TITLE OF INVENTION: METHODS OF IDENTIFYING FUNCTIONAL ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (1)..(9)
OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by T
OTHER INFORMATION: ell epitopes
                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                           OTHER INFORMATION: ER localization signal
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ZHOU, Chen
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88.9%;
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100.0%; Pred. No. 1.4e+05;
tive 0; Mismatches 0;
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Pred. No. 1.4e
0; Mismatches
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RESULT 6
US-11-225-686-211
; Sequence 211, Application US/112; Publication No. US20060019349A1
; Publication No. US20060019349A1
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Best Local S
Matches 4
GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/225,686
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SEQ ID NO 208
LENGTH: 7
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APPLICANT: LIU, Qia
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Publication No.
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APPLICANT: Collingwood, Trevor
TITLE OF INTENTION: Treatment of Neuropathic Pain with Zinc Finger Proteins
FILE REFERENCE: 019496-008220US
CURRENT FILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 60/560,535
PRIOR PRIOR PRIOR DATE: 2004-04-08
PRIOR PILING DATE: 2004-04-08
PRIOR PILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: US 60/576,757
PRIOR PILING DATE: 2004-06-02
NUMBER OF SEQ ID NOS: 227
NUMBER OF SEQ ID NOS: 227
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TYPE: PRT
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US-11-225-686-216
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US-11-225-686-216
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                                                       GENERAL INFORMATION:

APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLECTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT FILING DATE: 2005-08-11
PRIOR APPLICATION NUMBER: US/11/202,009
CURRENT FILING DATE: 2005-08-11
PRIOR APPLICATION NUMBER: US/9/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOCTWARE: PATENTIN Ver. 2.0
SEQ ID NO 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT FILING DATE: 2005-09-12
PRIOR APPLICATION NUMBER: US/11/225,686
CURRENT FILING DATE: 2005-09-12
PRIOR APPLICATION NUMBER: US/99/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 216
                                                                                                                                                                                                                                                                                                    Sequence 208, Application US/11202009 Publication No. US20060024726A1
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Best Local Similarity
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Publication No. US20060019349A1
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PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
LENGTH: 7
TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
PEATURE:
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Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
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; OTHER INFORMATION: Description of Artificial Sequence: example
US-11-202-009-208
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US-11-202-009-211
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TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT FILING DATE: 2005-08-11/202,009
CURRENT FILING DATE: 2005-08-11
PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR RILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN VEY. 2.0
SEQ ID NO 216
                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 216, Application US/11202009
Publication No. US20060024726A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 4; Conserv
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SEQ ID NO 211
LENGTH: 7
TYPE: PRT
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Publication No. US20060024726A1
PUBLICATION:
                                 Matches
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/202,009
CURRENT FILING DATE: 2005-08-11
PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR APPLICATION NUMBER: US/09/989,994
NUMBER OF SEQ ID NOS: 4085
NUMBER OF SEQ ID NOS: 4085
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                                                                                                                                                                             LENGTH:
                                                                                                                  FEATURE: OTHER INFORMATION: Description of Artificial Sequence: example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE: OTHER INFORMATION: Description of Artificial Sequence: example ZFP
                                                                                                                                               ORGANISM: Artificial Sequence
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3 RDDLLE 8
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                                 Conservative
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66.7%;
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                               Score 21; DB 7;
Pred. No. 1.4e+05;
1; Mismatches 1
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 21; DB 7;
Pred. No. 1.4e+05;
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                                                                Length 7;
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US-10-989-767A-219

Sequence 219, Application US/10989767A

US20060018917A1

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APPLICANY: ARM, CALLANA
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: UNCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: UNEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 511582005004
CURRENT APPLICATION NUMBER: US/10/989,767A
CURRENT APPLICATION NUMBER: 100/277,292
PRIOR APPLICATION NUMBER: 010/277,292
PRIOR FILING DATE: 2001-00-21
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR FILING DATE: 2001-08-20
PRIOR
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Best Local Similarity
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APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: OTHER CANCERS
FILE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 511582005004
CURRENT APPLICATION NUMBER: US/10/989,767A
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR FILING DATE: 2002-10-21
PRIOR PELICATION NUMBER: 09/935,430
PRIOR PELICAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: FARIS, MARY
APPLICANT: HUBERT, REN
APPLICANT: RAITANO, AF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE: OTHER INFORMATION: Chemically synthesized peptide motif
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RAITANO, ARTHUR
AFAR, DANIEL
LEVIN, ELANA
CHALLITA-EID, PI
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RAITANO, ARTHUR
AFAR, DANIEL
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Pred. No. 1.4e+05;
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Sequence 268, Application US/10989767A
Publication No. US20060018917A1
GENERAL INFORMATION:
APPLICANT: FARIS, MARY
APPLICANT: HUBERT, RENE

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APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 51158205004
CURRENT APPLICATION NUMBER: US/10/989,767A
CURRENT FILING DATE: 2004-11-15
CURRENT FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2000-08-22
PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR APPLICATION NUMBER: 60/282,739
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RESULT 14
US-10-989-767A-268
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PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 565
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 159
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                                                                                                                                                             Matches
                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                Query Match
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APPLICANT: HUBERT, REI
APPLICANT: RAITANO, AI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE: OTHER INFORMATION: Chemically synthesized peptide motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                  LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
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3; Conservative
                                                                                1 ILNDNAIES 9
                                                                                                                      1 VLRDDLLEA 9
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                                                                                                                                                               Conservative
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37.5%;
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Pred. No. 1.4e+05;
                                                                                                                                                                                    Score 21; DB 6;
Pred. No. 1.7e+02;
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RAITANO, ARTHUR

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APPLICANT: RAITANO, ARTHUR
APPLICANT: RAITANO, ARTHUR
APPLICANT: LEVIN, ELANA
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: O'SEPUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: O'STER CANCERS
FILE REFERENCE: 511582005004
CURRENT APPLICATION NUMBER: US/10/989,767A
CURRENT APPLICATION NUMBER: 10/277,292
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR FILING DATE: 2002-10-21
PRIOR PILING DATE: 2001-08-12
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,39
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 272
LENGTH: 10
TYPE: PRI
ORGANISM: Artificial Sequence
PRIOR INFORMATION: Chemically synthesized peptide motif
US-10-989-767A-272
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CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR FILING DATE: 2002-10-21
PRIOR PFLICATION NUMBER: 09/935,430
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR PFLICING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR FILING DATE: 2000-08-22
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR PILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 268
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US-10-989-767A-272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 50.0%; Score 21; DB 6; Length 10; Best Local Similarity 37.5%; Pred. No. 1.7e+02; Matches 3; Conservative 4; Mismatches 1; Indels
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TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 511582005004
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Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.

Minor histocompatibility antigen HA-1 T-cell epitope

#1

20-MAY-1999 AAW99195;

(first entry)

AAW99195 standard; peptide; 9 AA.

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ALIGNMENTS

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RESULT 1
AAW99195
ID AAW9
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Synthetic.
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Claim 1; Page 32; 47pp; English.

immune n bone

The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompactibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the

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RESULT 3
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                                                                                                                                                                                                                                                                                                     A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.
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 AAW99197 standard; peptide; 9 AA.
                                                                                                                                                   Sequence 9
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100.0%; Pr
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RESULT 4
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                                                                                          T-cell epitope; minor histocompatibility antigen transplant rejection; Graft-versus-Host Disease;
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                                                                                transplant rejection; Graineoplastic haematopoietic
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                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                      VLXDDLLEA
                                                                                                                                                                                                                                                                VLRDDLLEA
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                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                       (first entry)
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                                                                                                                               from the minor histocompatibility antigen HA-1.
                                  Location/Qualifiers
            /note=
                                                                                                                                                                                                                                                                                       9
                                                                                                                                                                                                      peptide;
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                                                                                                                                                                                                                                                                                                                          97.3%;
88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           47pp; English.
            "His or Arg"
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                                                                                  cell.
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                                                                                                                                                                                                                                                                                                                          Score 36;
Pred. No.
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                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                          DB 2;
2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HA-1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful to treat immune graft disease in bone
                                                                                                                                                                                                                                                                                                                                     Length
                                                                                               autoimmune disease;
                                                                                                          HA-1; vaccine;
                                                                                                                                                                                                                                                                                                                Indels
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WO9905173-A1

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RESULT 5
AAW97375
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents an immunogenic peptide constituting a T-cell epicope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graft-versus-Host Disease, or to treat autoimmune diseases.

Neoplastic haematopoietic cells presenting the peptides, in an HIA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B-and T-cells
Typing minor histocompatibility antigen HA-1 -
                      WPI; 1999-142960/12
                                                                                         23-JUL-1997;
02-JUN-1998;
                                                                                                                          23-JUL-1998;
                                                                                                                                                 04-FEB-1999
                                                                                                                                                                      WO9905313-A2
                                                                                                                                                                                              Homo
                                                                                                                                                                                                                 Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                HA-1 H-allele sequence
                                                                                                                                                                                                                                                                                         13-MAY-1999
                                                                                                                                                                                                                                                                                                                                     AAW97375
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Goulmy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   versus-host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-142855/12.
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                                                                     (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunogenic peptide from minor histocompatibility antigen HA-1 - for inducing tolerance to transplants and prevent rejection or {\sf gr}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-JUL-1998;
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les 9; Conser
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                                                                                                                                                                                                                                                                                                                                    standard;
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                                                                                         97EP-00202303.
98EP-00870125.
                                                                                                                          98WO-EP004928
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Pred. No.
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2e+06;
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by amplifying
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RESULT 6
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Best Local
The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used to screen for the HA-1 alleles. The HA-1 peptides can be used
                                                                                                                                                           Claim 13;
                                                                                                                                                                                         of genetic
                                                                                                                                                                                                                                                       WPI; 1999-142960/12
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02-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 9 AA;
                                                                                                                                                                                                      Typing minor histocompatibility antigen HA-1 - by amplifying identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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                                                                                                                                                                                                                                                                                                                    (UYLE-) RIJKSUNIV LEIDEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    identifying {\rm HA-1}\ R and {\rm HA-1}\ H alleles of {\rm HA-1}, used for, e.g. detection of genetic aberrances.
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                                                                                                                                                      English.
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Pred. No. 2e+0
0; Mismatches
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                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9 AA;
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                                                                                          fragment encoded by DNA sequence having SNP, creating variant epitopes, selecting epitopes binding to MHC protein. A protein of the invention has cytostatic activity, and may have a use in a vaccine. The method is useful for generating a SNP profile of one or more individuals from a given species by applying the method for several protein from the individuals, where the SNP profile was related to disease, preferably cancer. This is useful for diagnosing a disease in an individual by generating the SNP-related polymorphic profile. A method of the invention is useful for transplanting haematopoietic stem cells from a donor to a recipient and treating cancer, preferably leukaemia, and for determining the progression, regression or onset of a treated disease. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                               The invention relates to a novel method for providing epitopes of allelic variants of antigenic proteins from specific species based on single nucleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein, identifying, selecting allelic peptide/protein variants, expression product or its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Strittmatter W, Moll H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-JUN-2002; 2002EP-00013423
                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 82; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                          providing allelic variant epitope of protein based on single nucleotide
polymorphism by defining target protein, screening database of protein,
identifying, selecting allelic variant protein, creating variant
                                                                Sequence
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                 Local Similarity
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88.9%;
                   97.3%;
88.9%;
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    Score 36; DB
Pred. No. 2e+0
0; Mismatches
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2e+06;
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2e+06;
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel method for providing epitopes of allelic convariants of antigenic proteins from specific species based on single convicted polymorphism (SNP), by defining target protein/peptide or its converted protein graphism (SNP), by defining target protein/peptide or its converted protein graphism (SNP), by defining target protein/peptide or its converted protein graphism of the invention for selecting allelic peptide/protein variants, expression product or its converted protein graphism of the invention has converted protein graphism of the invention has converted protein for generating a SNP profile of one or more individuals from a converted protein graphism of the supplying the method for several protein from the converted protein graphism is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer. This is useful for diagnosing a disease in an individual by cancer in the SNP-related polymorphic profile. A method of the invention converted to transplanting cancer, preferably leukaemia, and for determining cancer, preferably leukaemia, and for determining cancer, preferably leukaemia, and for determining cancer is used in the exemplification of the invention.
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                                                                                                                                                                      Query Match
Best Local
                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                             Sequence 9 AA;
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                                                                                                                                     Local Similarity
nes 8; Conserv
                                                                  1 VLXDDLLEA
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VLRDDLLEA
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88.9%;
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                                                                                                                                                                             Score 36;
Pred. No.
                                                                                                                                            Mismatches
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ADT73023 ID ADT

ADT73023 standard; peptide; 9 AA.

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RESULT 9

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                                                                                                                           RESULT 10
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CC Poptide sequence is a human RSV high binding affinity peptide of the CC invention.
                                                                                                                                                                                                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                         Matches
SARS coronavirus spike protein HLA A 0201 T-cell epitope, SEQ:8174.
                                 16-JUN-2005
                                                                ABY01546;
                                                                                            ABY01546 standard;
                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-APR-2003; 2003EP-00447095
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             human respiratory syncytial virus; cytotoxic T lymphocyte; CTL; MHC epitope; DNA-based immunisation; RSV infection; virucidal; vaccine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human RSV L high affinity binding peptide SeqID 919
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                                                                                                                                                                                                                                     Local Similarity
nes 4; Conserv
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                             (first entry)
                                                                                            peptide;
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Pred. No. 2e+0
3; Mismatches
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22-APR-2003;
22-APR-2003;
23-APR-2003;
24-APR-2003;
24-APR-2003;
05-MAY-2003;
14-AUG-2003;
                                                                                         23-SEP-2003;
11-OCT-2003;
                                                                                                                                                                                                                                                                    18-APR-2003;
                                                                                                                                                                                                                                                                                                        15-APR-2003;
16-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                12-APR-2003;
13-APR-2003;
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11-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SARS coronavirus.
                                                                                                                                                                                                                                                                                      17-APR-2003
                                                                                                                                                                                                                                                                                                                                              14-APR-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO2004092360-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vaccine; nucleic acid vaccine; drug screening; diagnosis; SARS coronavirus infection; infection; respiratory disease; virucide.
(CHIR ) CHIRON CORP.
                               2003US-046483BP.

2003US-0464899P.

2003US-0465273P.

2003US-046535P.

2003US-0473144P.

2003US-0473144P.

2003US-0473149.

2003US-0473149.

2003US-0473149.

2003US-0473149.

2003US-0473149.

2003US-0473149.

2003US-0505652P.

2003US-0510781P.

2004US-0536177P.

2004US-0536177P.
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2003US-046274BP.
2003US-046310pp.
2003US-0463460P.
2003US-046386BP.
2003US-0463983P.
2003US-0463983P.
2003US-0463983P.
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2003US-0462465P
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Rappuoli R, Masignani V, Stadler K, Gregersen J, Chien D, Han Polo J, Weiner A, Houghton M, Song HC, Seo MY, Donnelly JJ; Klenk HD, Valiante N;

PI; 2004-766863/75

Novel isolated polypeptide e.g. spike polypeptide, Env polypeptide, of severe acute respiratory syndrome virus (SARS), useful as vaccine for SARS.

Disclosure; SEQ ID NO 8174; 839pp; English.

The invention relates to isolated polypeptides of the severe acute CC respiratory syndrome (SARS) corconavirus. The polypeptides include spike CC (S or E2), env (E or sM), membrane (M or E1), hemagglutinin-esterase (HE CC or E3), and nucleocappid (N) polypeptides, and the ORF1a and ORF1ab CC (replicase) polypeptides and their proteolytic fragments. The invention CC acids encoding the SARS virus polypeptides, primers specific for SARS (Virus nucleic acid sequences; kits for amplifying SARS virus target CC virus nucleic acids; a double-stranded RNA molecule 10-30 nucleotides in length CC which is able to inactivate the SARS virus in a mammalian cell; an CC expression construct for recombinant expression of a SARS virus spike CC viral antigen. The invention additionally provides a vaccine for the CC viral antigen. The invention additionally provides a vaccine for the CC wistled SARS virus, an attenuated SARS virus, a split SARS virus, a cc virus antigens, an attenuated SARS virus, a split SARS virus and comprising one or more SARS virus and one or more comprising one or more SARS virus antigens; methods of comprising one or more SARS virus antigens and one or more comprising one or more SARS virus antigens and one or more comprising one or more SARS virus antigens and one or more comprising one or more SARS virus antigens and one or more comprising one or more SARS virus antigens and one or more comprising one or more SARS virus antigens and one or more comprising one or more SARS virus antigens and one or more comprising one or more SARS virus antigens and one or more comprised to a SARS patient comprised condition of the comprised conditions. The sars virus polypeptides and collected acids can be used in the preparation and manufacture of vaccines conditions and manufacture of vaccines conditions.

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RESULT 11
The invention relates to a cytotoxic Yersinia pestis T-cell epitope, competition and of the 1000 sequences of 9 amino acids appearing as ADZ49933-ADZ50932. Also included are predicting peptides that are ceptitopes or can be used as diagnostic tools (comprising predicting which peptides bind to a HMC molecule (not defined) with high affinity using a neural network with at least one of the following features: some or all cof the inputs to the neural networks are generated using a hidden Markov model; or some or all of the inputs are encoded by an amino acid substitution matrix, different from an identity matrix) and a vaccine or diagnostic tool using a limited number such as at least 1, 2, 3, 4, 5, 8, 16, 32, 64, 128, 256, 512 of the peptides of ADZ49933-ADZ50932. In predicting peptides, the prediction of the neural network is combined with prediction or measurement of one of the following: proteasomal cleavage sites; MHC binding; presence of sequence or related sequence(s) in patent databases; TAP binding; gene or protein expression level; function of the protein; localization of the protein; and similarity to self proteins. The epitope is useful as a vaccine or diagnostic tool, and
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Best Local
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                                                                                                                                                                                                                                                                                                                                                      in a
                                                                                                                                                                                                                                                                                                                                                                     New cytotoxic Yersinia pestis T-cell epitope comprising useful as a vaccine or diagnostic tool and for inducing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-APR-2005
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                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 993; 235pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pestis infection; antibacterial; infection; biological warfare.
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RESULT 12
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Matches
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Best Local
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                                             Sequences AAU71093-AAU71796 represent the 1039388-related protein and peptide fragments of the protein. 1039388 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 1039388 polynucleotide, its related protein and peptide fragments and specific PCR primers are therefore useful for diagnosing and treating cancer. A vector comprising a polynucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 1039388 related protein, and a ribozyme capable of cleaving a polynucleotide having the 1039388 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 1039388. The sequences can be used in diagnostic methods to monitor the level of 1039389 gene products in serum, blood, urine and tissue and to thereby detect the presence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tumour; cancer; cytostatic; gene therapy; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine; tissue; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromosome 9q13-q21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human MHC class I molecule HLA-A2 binding 103P3E8 peptide #18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU71211 standard; peptide; 9
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                                                                                                                                                                                                                                                                                                      Disclosure; Page 84; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                     Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e. prostate cancer, comprises determining presence of aberrant 103P3E8 ge
                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-061976/08.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Faris M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-APR-2000; 2000US-0196647P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-OCT-2001
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                                 cells
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62.5%;
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Pred.
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                                                                                                                                                                  Sequences AAU71093-AAU71796 represent the 103P3E8-related protein and peptide fragments of the protein. 103P3E8 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expressed in many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 103P3E8 polymuclectide, its related protein and peptide fragments and specific PCR primers are therefore useful for diagnosing and treating cancer. A vector comprising a polymuclectide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 103P3E8-related protein, and a ribozyme capable of cleaving a polymuclectide having the 103P3E8 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 103P3E8. The sequences can be used in diagnostic methods to monitor the level of 103P3E8 gene products in serum, blood, urine and tissue and to thereby detect the presence of
                                                                                                                           Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 90; 128pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e.g. prostate cancer, comprises determining presence of aberrant 103P3E8 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-061976/08.
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ADT72767 standard; peptide; 9 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 16; SEQ ID NO 1586; 143pp; English.
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Matches 4
                                                                                                                                                                                                              Sequence 9 AA;
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MHC epitope; DNA-based immunisation; RSV infection; virucidal; vaccine.
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                                                                     hucolin, 75K chain hypothetical prote elastase - Pseudom protein kinase C i trypsin (EC 3.4.21 T-cell receptor be macrophage cytotox biotin A - Citroba T-cell receptor be cerebellar degener T-cell receptor be acetylcholinestera calsequestrin, fas T-cell receptor be glycine cleavage s phosphoglycerate t pilE protein - Esc chittin-binding protein cleavage s phosphoglycerate t cile protein - Esc chittin-binding protein - Esc
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4 DDL 6 ||| 2 DDL 4

Best Local Matches	RESULT 2 PC4131 hypothetical protein C;Species: Pseudomona C;Date: 20-Apr-2000 # C;Accession: PC4131 R;Kawasaki, S.; Arai, Gene 167, 87-91, 1995 A;Title: Sequencing a y for biosynthesis of A;Reference number: J A;Accession: PC4131 A;Status: preliminary A;Molecule type: DNA A;Residues: 1-8 <kaw- a;cross-references:="" a;note:="" c;superfamily:="" is="" n="" orf="" pseudo<="" th="" this="" u=""><th>Query Match Best Local; Matches Qy 4 Db 4</th><th>RESULT 1 S68004 hucolin, 75K C;Species: Ho C;Date: 15-Fe C;Accession: R;Edgar, P.F. REBS Lett. 37 A;Title: Huco A;Reference n A;Accession: A;Cacss-refer</th><th></th><th>0 1 2 8 4 8 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4</th></kaw->	Query Match Best Local; Matches Qy 4 Db 4	RESULT 1 S68004 hucolin, 75K C;Species: Ho C;Date: 15-Fe C;Accession: R;Edgar, P.F. REBS Lett. 37 A;Title: Huco A;Reference n A;Accession: A;Cacss-refer		0 1 2 8 4 8 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4
Similarity 3; Conservat	rotein 8 (udomonas a -2000 #seq -2111 #seq -2111 H. 1, 195 1, 195 ancing and ncing and ncing of he mber: JC45 C4131 indinary indinary e: DNA 8 <kaw- <kaw-="" b="KAW-" b<="" td=""><td>h Similarity 3; Conservat 4 DDL 6 4 DDL 6</td><td>chain - hun mo sapiens b-1997 #sec \$68004 5, 159-161, lin, a new umber: \$68004 liminary pe: proteir -7 <edg> ences: UNIE</edg></td><td></td><td>11 29.7 10 27.0 10 27.0</td></kaw->	h Similarity 3; Conservat 4 DDL 6 4 DDL 6	chain - hun mo sapiens b-1997 #sec \$68004 5, 159-161, lin, a new umber: \$68004 liminary pe: proteir -7 <edg> ences: UNIE</edg>		11 29.7 10 27.0 10 27.0
op.	[imported] - aeruginosa quence_revis 1.; Igarashi, 1 characteriz eme dl. 1552; MUID:96 1552; MUID:96 178412; annotated i nas stutzeri 1784 72%; S	43.2%; 100.0%; ;ive	n (fragment) man) ence_revisic ence_revisic orticosteroi 4; MUID:9608		タタ4466777778888833333333333333333333333333
Pred. No. 2.8e+05 0; Mismatches 0	Pseudomonas aerugino on 20-Apr-2000 #text Y.; Kodama, T. Lion of the downstre Lion of the downstre PMID:8566817 44254; PMID:8566817 44254; PMID:8566817	Score 16; DB 2; Length 7; Pred. No. 2.8e+05; 0; Mismatches 0; Indels	13-Mar-1997 #text	ALIGNMENTS	A12872 A61386 A48360 A26209 A26209 I148394 I148394 I148394 I148394 I148394 S78024 S78024 S78024 S78029 S29735 S29735 PC2370 PC23
els 0; Gaps 0;	» та	lels 0; Gaps 0;	_change 17-Mar-1999 rom human plasma with structural		transaldolase (BC macrophage inhibit gamma subunit of p protein-glutamine glycoprotein compo alpha-tubulin - Ch ribosomal protein formylglycinamide platelet glycoprot polyphosphate-gluc probable H+-transp NADH2 dehydrogenas peptidyl-dipeptida hypothetical prote Ig heavy chain CRD fibroblast growth

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C;Accession: S20446
R;Kessler, E.; Safrin, M.; Peretz, M.; Burstein, Y.
FEBS Lett. 299, 291-293, 1992
A;Title: Identification of cleavage sites involved in proteolytic processing of Pseudomc A;Reference number: S20446; MUID:92183956; PMID:1544509
A;Accession: S20446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Description: Purification A; Reference number: A59489 A; Accession: A59489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein kinase C inhibitor - rat (fragment) C;Species: Rattus norvegicus C;Date: 25-Aug-2003 #sequence_revision 25-AC;Accession: A59489
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C;Species: Pseudomonas aeruginosa
C;Date: 19-Mar-1997 #sequence_revision 24-Jul-1997 #text_change 24-Jul-1997
                                                                               A;Cross-references: UNIPROT:Q7M390; UNIPARC:UPI000017C43F
C;Keywords: hydrolase; protein digestion; serine proteinase; zymogen
F;1-8/Domain: activation peptide #status experimental <APT>
                                                                                                                                                                                                                           R;Bricteux-Gregoire, S.; Schyns, R.; Florkin, M.;
Biochim. Biophys. Acta 386, 244-255, 1975
A;Title: N-terminal amino acid sequence of trypsir
                                                                                                                                                                                                                                                                                               trypsin (EC 3.4.21.4) precursor - minke whale (fragment)
C;Species: Balaenoptera acutorostrata (minke whale, lesser rorqual)
C;Date: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  submitted to the Protein Sequence Database, June 2003 A; Description: Purification of PKCI from rat liver.
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A;Note: p-Hydroxyacetophenone-Sepharose
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           Score 13; DB 2;
Pred. No. 2.8e+05;
1; Mismatches 1
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A60427
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J. Exp. Med. 174, 1467-1476,
A;Title: Analysis of T cell;
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A; Residues: 1-9 < GOL>
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              A; Molecule type: DNA
A; Residues: 1-4 < RES
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macrophage cytotoxicity-inducing factor, 29K - human (fragment)
C.Species: Homo sapiens (man)
C.Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
C.Accession: A60427
R.Jones, C.M.; Prince, C.A.; Williams, J.S.
RayD. Hematol. 19, 704-709, 1991
A.Title: Purification and amino acid analysis of a human macrophage cytotoxicity-inducin A.Reference number: A60427; MUID:91372335; PMID:1909970
A.Reference A60427
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A;Acession: PH0942
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C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
C;Accession: PH9942
                                                     R;Shiuan, D.; Campbell, A.
Gene 67, 203-211, 1988
A;Title: Transcriptional regulation and gene
A;Reference number: I40697; MUID:89006280; PM
                                                                                                                                                         biotin A - Citrobacter freundii (fragment)
C;Species: Citrobacter freundii
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 09-Jul-2004
C;Accession: 140597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: protein
A;Residues: 1-9 <JON>
A;Residues: 1-9 <JON>
A;Cross-references: UNIPROT:Q7M4R5; UNIPARC:UPI0000142BE0
A;Note: the sequence from the text on page 706 is inconsistent with that from page 708
C;Keywords: cytokine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: UNIPARC:UPI000017C9DD A;Experimental source: complete Freund's A;Experiments source: the authors translated the codon' C;Keywords: T-cell receptor
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A;Status: preliminary; translated from GB/EMBL/DDBJ
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Pred. No.
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Pred. No. 2.8e+05;
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PMID:2971595
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R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0601
A;Accession: PT0601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T-cell receptor beta chain V-D-J region (120-2K) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 17-Jul-1992 #sequence revision 17-Jul-1992 #text_change 09-Jul-2004 C;Accession: PT0601; PT0617; PT0694
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PT0679
A;Cross-references: UNIPARC:UPI000017C7E6
A;Experimental source: day 18 fetal thymu:
C;Keywords: T-cell receptor
                                                                  A; Molecule type: DNA
A; Residues: 1-5 < FE2>
                                                                                                                                                                                       A;Status: translation not shown A;Molecule type: mRNA A;Residues: 1-5 <PE3>
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A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0679
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0679; PT0708
                                                                                                                                        A;Residues: 1-5 \ruse NIPARC:UPI000017C7E6
A;Cross-references: UNIPARC:UPI000017C7E6
                                                                                                                                                                                                                                                                          A;Cross-references: UNIPROT:Q64512; UNIPARC:UPI000017C7E6
A;Experimental source: newborn thymus, strain BALB/c, clone 120-2K
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A; Residues: 1-5 < FEE>
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A; Residues: 1-5 < FEE>
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A; Residues: 1-5 < FE2>
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A;Experimental source: day 18 fetal thymus,
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Experimental source: newborn thymus, strain BALB/c,
Keywords: T-cell receptor
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100.0%; Pred. No. 2.8e+05;
tive 0; Mismatches 0;
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                   strain BALB/c,
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es 0;
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acetylcholinesterase (EC 3.1.1.7) 5.6S form - Pacific electric ray (fragment) (;Species: Torpedo californica (Pacific electric ray)
C;Date: 16-Mar-1990 #sequence_revision 16-Mar-1990 #text_change 08-Nov-1996
C;Accession: A34026
R;Gibney, G.; MacPhee-Quigley, K.; Thompson, B.; Vedvick, T.; Low, M.G.; Taylor, S.S.; '1
J. Biol. Chem. 263, 1140-1145, 1988
A;Title: Divergence in primary structure between the molecular forms of acetylcholinests A;Reference number: A34026; MUID:88087239; PMID:3335534
A;Accession: A34026
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J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T ce
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A;Title: Cerebellar degeneration-related antigen: a highly conserved neuroectodermal ma A;Reference number: A35640; MUID:90222173; PMID:2326268
A;Accession: B35640
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A;Molecule type: mRNA
A;Residues: 1-6 <FEE>
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                                                                                                                                                                                  RESULT 13
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A; Residues: 1-6 < CHE>
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100.0%; Pr
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calsequestrin, fast skeletal muscle - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 23-Aug-1991 #sequence_revision 23-Aug-1991 #text_change 23-Feb-1997
C;Accession: B39040
R;Cala, S.E.; Jones, L.R.
J. Biol. Chem. 266, 391-398, 1991
A;Title: Phosphorylation of cardiac and skeletal muscle calsequestrin isoforms by casein A;Reference number: A39040; MUID:91093153; PMID:1985907
A;Accession: B39040
                                                                                                                                                                                                                                                                                                                                                                            T-cell receptor beta chain V-D-J region (111-1AG) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0628
R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
J. Exp. Med. 174, 115-124, 1991
A;Fitle: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0628
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A;Residues: 1-7 <GIB>
A;Cross-references: UNIPARC:UPI000017BF32
C;Keywords: alternative splicing; carboxylic ester hydrolase
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A;Molecule type: mRNA
A;Molecule: 1-7 <FEE>
A;Cross-references: UNIPARC:UPI000017C7CD
A;Experimental source: newborn thymus, strain BALB/c C;Keywords: T-cell receptor
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A;Molecule type: protein
A;Residues: 1-7 <CAL>
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Search completed: April 6, 2006, 16:26:10 Job time: 20.6667 secs
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Best Local S
Matches 2
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ilarity 100.0%; Pred. No. 2.8e+05;
Conservative 0; Mismatches 0;
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Maximum Match 100%
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Perfect score:
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1: uniprot_sprot:*
2: uniprot_trembl:*
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Q9FSZ2_CICAR
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Q8UTD7_9HIV1
Q99HD8_9SMEG
Q6UTB2_CRYNV
Q9HD84_ASPFL
Q7M390_BALAC
Q9HD84_ASPFL
Q7M390_BALAC
Q8H9K1_9CAUD
Q8SBJ0_BPR69
Q6JC68_SOVEN
P72279_RHOGO
Q51594_9ZZZZ
Q7M4R5_HUMAN
Q6UVK2_MALLO
Q56ST1_9UIPS
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P82568_STRPY
Q9XJN0_9VIRU
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53.811 Million cell updates/sec
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Q70sm2 homo sapien
Q81pt5 zea mays (m
P82568 streptococc
Q9xjn0 bacteriopha
Q9fg22 cicer ariet
Q9g2a8 mus musculu
Q85710 rous sarcom
Q8utd7 human immun
Q99182 gnatholebia
Q6u772 cryptococcu
Q99464 aspergillus
Q7m390 balaenopter
Q8h9k1 bacteriopha
Q6u568 glycine max
P7279 rhodococcus
Q51594 plasmid col
Q7m4r5 homo sapien
Q9bqr4 homo sapien
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RESULT 2 QBLPT5 MAIZE ID QBLPT5 MAIZE PRELIMINARY; PRT; 9 AA. QBLPT5; AC QBLPT5; DT 01-OCT-2002 (TrEMBLrel. 22, Created) DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update) DT 01-OCT-2002 (TrEMBLrel. 22, Last annotation update) DE Beta-expansin-like protein (Fragment). OS Zea mays (Maize). OS Zea mays (Maize). OC ENCATYOCL; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta OC Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea. OX NCBI TaxID=4577; RN [1] RP NUCLEOTIDE SEQUENCE. RA Ching A.S., Caldwell K.S., Jung M., Dolan M., Smith O.S., Tingey RA Morgante M., Rafalski J.A.; RL Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases. EMBL; AY094310; AAM21836.1; -; Genomic_DNA.	Query Match 43.2%; Score 16; DB 2; Length 9; Best Local Similarity 60.0%; Pred. No. 2.2e+06; Matches 3; Conservative 1; Mismatches 1; Indels 0; C Qy 5 DLLEA 9 : Db 3 DTMEA 7	NCBI_TaxID=9606; [1] NUCLEOTIDE SEQUENCE. TISSUE=Mammary gland; Pinte S., Guerardel C., Deltour S., Godwin A.K., Leprince D.; "Identification of a second G-C-rich promoter conserved in the murine and rat tumor suppressor genes HIC1."; Oncogene 23:4023-4031(2004). EMBL; AJ550616; CAD79467.1; -; mRNA. NON TER 9 SEQUENCE 9 AA; 964 MW; 5B5E6DDB16B1AAA7 CRC64;	RESULT 1 Q70SM2_HUMAN ID Q70SM2_HUMAN PRELIMINARY; PRT; 9 AA. AC Q70SM2; DT 05-JUL-2004 (TrEMBLrel. 27, Created) DT 05-JUL-2004 (TrEMBLrel. 27, Last sequence update) DT 05-JUL-2004 (TrEMBLrel. 27, Last annotation update) DT Hypermethylated in cancer 1 (Fragment). GN Name=HICL; OS Homo sapiens (Human). OC Enkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; OC Homo.	2 13 35.1 9 2 Q56SV1-9IPS Q56SV2 3 13 35.1 9 2 Q56SV2-9DIPS Q56SV2 4 13 35.1 9 2 Q56SV3-9DIPS Q56SV3 5 13 35.1 9 2 Q56SV4-9DIPS Q56SV4 5 13 35.1 9 2 Q56SV4-9DIPS Q56SV4 ALIGNMENTS	13 35.1 9 2 Q56SUI_9DIPS Q56su1 13 35.1 9 2 Q56SUZ_9DIPS Q56su2 13 35.1 9 2 Q56SUZ_9DIPS Q56su3 13 35.1 9 2 Q56SUJ_9DIPS Q56su4 13 35.1 9 2 Q56SUS_9DIPS Q56su5 13 35.1 9 2 Q56SUS_PDIPS Q56su6 15 35.1 9 2 Q56SUS_PDIPS Q56su6 16 35.1 9 2 Q56SUS_PDIPS Q56su6 17 35.1 9 2 Q56SUS_PDIPS Q56su6 18 35.1 9 2 Q56SUS_PDIPS Q56su6 19 35.1 9 2 Q56SUS_PDIPS Q56su6 19 35.1 9 2 Q56SUS_PDIPS Q56su6
eophyta; Tingey S.,	0; Gaps 0;	, U	ostomi; ; Hominidae;	viburnum viburnum viburnum viburnum	

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01-NOV-1999 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
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Q9XJNO;
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01-OCT-2000 (TrEMBLrel. 15, Last sequence)
01-OCT-2003 (TrEMBLrel. 25, Last annotat.
Unknown protein from 2D-page (Fragment).
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Viruses; dsRNA viruse
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VanBogelen R.A.;
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MEDLINE=99350412; Pu
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Hoogstraten D.;
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid
eurosids I; Fabales; Fabaceae; Papilionoideae; Cicereae; Cicer.
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01-MAR-2001 (TrEMBLrel. 16,
01-OCT-2003 (TrEMBLrel. 25,
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Q9QZA8;
01-MAY-2000 (TrEMBLrel. 13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dopico B., Jimenez T., Labrador E.; Submitted (SEP-2000) to the EMBL/GenBank/DDBJ EMBL; AJ299069; CAC10216.1; -; mRNA.
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09FSZ2;
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Submitted (OCT-1999) to the EMBL/GenBank/DDBJ
EMBL, AF192526; AAF04843.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Mus.
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Name=Clec2i; Synonyms=Dcl1;
Mus musculus (Mouse).
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, MGI:2136650; Dcll.
GO:0009897; C:external side of plasma membrane;
GO:0005887; C:external to plasma membrane; TAS.
GO:0005887; F:natural killer cell lectin-like re
GO:0005515; F:protein bänding; IPI.
GO:0016066; P:cellular defense response (sensu V
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NOVITSKY V.A., McLane M.F., Chigwedere P., Ndung'u T., Kl

Novitsky V.A., McLane M.F., Rybak N., Gaseitsiwe S.,

Chang S.-Y., Peter T., Thior I., Rybak N., Gaseitsiwe S.,

Marlink R., Lee T.-H., Essex M.;

Submitted (OCT-2001) to the EMBL/GenBank/DDBJ databases.

Submitted (OCT-2001) to the EMBL/GenBank DDBJ databases.

SEQUENCE 9 AA; 1102 MW; 188BD40B17272440 CRC64;
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01-MAR-2002
01-OCT-2002
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DOI=10.1128/JVI.76.11.5435-5451.2002;
Novitsky V., Smith U.R., Gilbert P., McLane M.F.,
Williamson C., Ndung'u T., Klein I., Chang S.-Y.,
Foley B.T., Gaolekwe S., Rybak N., Gaseitsiwe S.,
Marlink R., Lee T.-H., Essex M.;
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MEDLINE-84115080; PubMed-6319754;
MEDLINE-84115080; PubMed-6319754;
Lerner T.L., Hanafusa H.;
"DNA sequence of the Bryan high-titer strain of Rous sarcoma virus:
"DNA sequence of the Bryan high-titer strain of Rous sarcoma virus:
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"DNA sequence of the Bryan 
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Primate lentivirus group.
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RESULT 9
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099182;
DT 01-MAY-1999 (T
DT 01-MAY-1999 (T
DT 01-MAY-1999 (T
DT 01-UN-2003 (T
DE CYCCCHOME OXI.
GN Name=COI;
OS Gnatholebias z
OG Mitochondrion.
OC Eukaryota; Met
OC Actinopterygii
OC Actinopterygii
OC Actinopterygii
OC Actinopterygii
OC Actinopterygii
OC Actinopterygii
OC ACTINOCONTIO
OX NCBI TAXID=135
RP NUCLEOTIDE SEQ
MEDLINE=200729
RA Murphy W.J., T
RT "Phylogeny of
RT "Phylogeny of
RT "Phylogeny of
RT GCyprinodontif
RT Sequences.";
RL Mol. Phylogene
DR EMBL; AF002591
DR GO; GO:0005739
KW Mitochondrion.
FT NON TER
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QEUTR2 CRYNV
ID QEUTR2 CRYNV
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DT 05-UUL-2
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DT 05-UUL-2
DT 05-UUL-2
CRYPTCOCO
OC ENEARYOUT
OC Tremello
OX NCBI Tax
RN NUCLEOTI
RC STRALN=H
RA STUART L
RL SUBMITTE
DR EMBL; AY
DR GO; GO:0
KW MGNOOXY9
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Best Local
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QGU7R2;
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
                                                                                                                                                                   Stuart L.T., Allen A., Dietrich F.S.; Submitted (AUG-2003) to the EMBL/GenBank/DDBJ databases EMBL; AY376718; AAQ88127.1; -; mRNA. GO; GO:0004506; F:squalene monooxygenase activity; IEA. Monooxygenase; Oxidoreductase.
                                                                                                                                                                                                                                                                                                                          Cryptococcus neoformans var. grubii H99.
Eukaryota; Fungi; Basidiomycota; Hymenom
Tremellomycetidae; Tremellales; Tremella
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05-JUL-2004 (TrEMBLrel. 27, Last seque)
05-JUL-2004 (TrEMBLrel. 27, Last anno Squalem monooxygenase (EC 1.14.99.7)
                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                             STRAIN=H99
                                                                                                                                                                                                                                                                                                          NCBI_TaxID=235443;
                                                                                                                                                                                                                                                                                                                                                                                  Name=ERG1;
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NCBI_TaxID=135316;
                                                                                                                                                                                                                                                                           NUCLEOTIDE SEQUENCE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-20072928; PubMed-10603257; DOI-10.1006/mpev.1999.0656; Muzphy W.J., Thomerson J.E., Collier G.E.; Phylogeny of the Neotropical killifish family Rivulidae (Cyprinodontiformes, Aplocheiloidei) inferred from mitochondri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mitochondrion.
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                                                                  Conservative
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                                                                                                                                      811 MW;
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40.0%;
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Pred. No. 2.2e+06;
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RESULT 12
Q7M390 BALAC PRELIMINARY;
AC Q7M390;
AC Q7M39
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ID Q9HDS4 ASI
ID Q9HDS6
AC Q9HDS
DT 01-MJ
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DT 01-MJ
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01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2001 (TrEMBLrel. 16,
TrpC polyprotein (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAR-2004 (TrEMBLrel. 26, Created)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last sequence)
11-MAR-2004 (TrEMBLrel. 26, Last sequence)
12-Marcological (EC 3.4.21.4) precursor (Fragment).
13-Malenoptera acucorostrata (Minke whale) (Lesser rorqual).
14-Bulaeroptera acucorostrata (Minke whale) (Lesser rorqual).
15-Mammalla; Eutheria; Chordata; Craniata; Vertebrata; Euteleostomi;
16-Mammalla; Eutheria; Laurasiatheria; Cetartiodactyla; Cetacea;
16-Mammalla; Balaenopteridae; Balaenoptera.
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Eurotiales; Trichocomaceae; mitosporic
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Q9HDS4;
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                                                                                                                                                                                 rorqual, Balaenoptera acutorostrata (Cetacea). Simultaneous isolat of trypsinogen, chymotrypsinogen and ribonuclease from pancreas."; Biochim. Biophys. Acta 386:244-255(1975).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "N-terminal amino acid sequence of trypsinogen from the rorqual, Balaenoptera acutorostrata (Cetacea). Simultar of trypsinogen, chymotrypsinogen and ribonuclease from Biochim. Biophys. Acta 386:244-255(1975).
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itosporic Trichocomaceae; Asperg
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RESULT 13
QRH9KI 9CAUD PRELIMINARY;
ID QRH9KI;
AC QRH9KI;
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DT 01-MAR-2003 (TrEMBLrel. 23,
DE GP30.2 protein (Fragment).
GN Name-30.2;
GN Name-30.2;
OS Bacteriophage LZ10.
CViruses; dabNA viruses, no
CC T4-like viruses,
CX NCBI TaxID-192973;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RA KOlesinskiene G., Nivinskas
RL Submitted (APR-2002) to the
DR EMBL; AJ458400; CAD30256.1;
FT NON TER
SQ SEQUENCE 8 AA; 918 MW;
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Q8SBJ0_BPR69
ID Q8SBJ0_B
RESULT 15
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T4-like viruses.
NCBI_TaxID=12353;
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1: /cgm2_6/ptodata/1,
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                                      ; OTHER INFORMATION: US-09-269-250E-20
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Query Match

97.3%;

Score

36; BB 2;

Length

Description

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Artificial Sequence:

Exon

Fragments

RESULT 2 US-09-269-250E-20 US-09-269-250E-20 Sequence 20, Application US/09269250E Patent No. 6830883 CENERAL INFORMATION: APPLICANT: Goulmy, Elsa TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1 FILE REFERENCE: 58994 CURRENT APPLICATION NUMBER: US/09/269,250E CURRENT FILLING DATE: 1999-05-21 NUMBER OF SEQ ID NOS: 38 SOFTWARE: Patentin version 3.1 SEQ ID NO 20 LENGTH: 9 TYPE: PRT ORGANISM: Artificial Sequence FEATURE: ORGANISM: Artificial Sequence	Query Match 97.3%; Score 36; DB 2; Length 9; Best Local Similarity 88.9%; Pred. No. 4.6e+05; Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0; Qy 1 VLXDDLLEA 9	RESULT 1 US-09-269-250B-18 ; Sequence 18, Application US/09269250E ; Patcent No. 6830883 ; GENERAL INFORMATION: APPLICANT: Goulmy, Elsa ; TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-: FILE REFERENCE: 58994 ; CURRENT APPLICATION NUMBER: US/09/269,250E ; CURRENT FILING DATE: 1999-05-21 ; NUMBER OF SEQ ID NOS: 38 ; SOFTWARE: Patcentin version 3.1 ; SEQ ID NO 18 ; LENGTH: 9 ; TYPE: PRT ; ORGANISM: Artificial Sequence ; PEATURE: ; OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments US-09-269-250E-18	28 21 56.8 9 2 US-09-217-609A-11 Sequence 11, Appl 29 21 56.8 9 2 US-08-873-235B-11 Sequence 11, Appl 30 21 56.8 9 2 US-09-586-472-72 Sequence 72, Appl 31 21 56.8 9 2 US-09-586-472-78 Sequence 78, Appl 32 21 56.8 9 2 US-09-528-706-78 Sequence 78, Appl 33 21 56.8 9 2 US-09-528-706-78 Sequence 78, Appl 33 21 56.8 9 2 US-09-528-706-78 Sequence 78, Appl 34 21 56.8 9 2 US-09-528-706-78 Sequence 78, Appl 35 20 54.1 6 2 US-09-808-126-11 Sequence 11, Appl 36 20 54.1 6 2 US-09-808-126-11 Sequence 11, Appl 39 20 54.1 8 2 US-09-803-951-11 Sequence 11, Appl 40 20 54.1 8 2 US-08-747-599A-50 Sequence 11, Appl 51.4 6 2 US-08-747-599A-50 Sequence 51, Appl 42 19 51.4 6 2 US-08-74-599A-51 Sequence 51, Appl 51.4 6 2 US-08-74-599A-51 Sequence 373, Appl 51.4 6 2 US-08-74-599A-51 Sequence 373, Appl 51.4 6 2 US-08-974-549A-373 Sequence 373, Appl 51.4 6 2 US-09-402-181B-373 Sequence 373, Appl 51.4 6 2 US-09-402-181B-373 Sequence 373, Appl 51.4 6 2 US-09-721-456-373 Sequence 373, Appl 51.4 6 2 US-0
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Best Local Matches

Similarity 88.9%; 8; Conservative

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Pred. No. 4.6e+05; 0; Mismatches 1;

Indels

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Gaps

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US-09-269-250E-29

US/09269250E

GENERAL INFORMATION: Sequence 29, Application Patent No. 6830883

SOFTWARE: PatentIn version 3.1 SEQ ID NO 29

ENGTH:

CURRENT APPLICATION NUMBER: US/09/269,250E CURRENT FILING DATE: 1999-05-21 NUMBER OF SEQ ID NOS: 38

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GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
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                                                                                     Query Match
Best Local
                                                                 Matches
                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Histocompatibility antigen FEATURE:
                                                                                                                                                                    NAME/KEY: PEPTIDE LOCATION: (3)
LOCATION: (3)
OTHER INFORMATION: AMINO ACID X REPRESENTS A HISTIDINE OR AN ARGININE OTHER INFORMATION: RESIDUE
                                                                                                                                                                                                                                                                                                                      LENGTH:
1 VLXDDLLEA 9
                                                             h 97.3%; Score 36; DB Similarity 100.0%; Pred. No. 4. 9; Conservative 0; Mismatches
                                                                                       DB 2; Le
                                                                                                           Length 9;
                                                                        Indels
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                                                                        Gaps
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APPLICANT: Goulmy, Elsa

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APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           residue
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Sequence 38, Application Patent No. 6830883 GENERAL INFORMATION:
                                                   RESULT 7
US-09-269-250E-38
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                                                                                                                                                                                                                                                        US-09-489-760-5
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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                      Query Match
Best Local S
Matches 8
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Best Local (
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                                                                                                                                                                                                                                                                                                                         SEQ ID NO 5
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CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: KIAA0223 partial complementary DNA
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Pred. No. 4.6e+05;
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RESULT 4 US-09-489-760-1

Sequence 1, Application US/09489760 Patent No. 6878375

US-09-269-250E-29

TYPE: PRT
ORGANISM: Human
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a histidine

Matches Query Match

Local Similarity les 9; Conserv

Conservative

97.3%; Score 36; DB 100.0%; Pred. No. 4.0 ative 0; Mismatches

DB 2, 3. 4.6e+05; 0;

Indels

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Gaps

Length 9;

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1 VLXDDLLEA 9

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; LENGTH: 8
; TYPE: PRT
; ORGANISM: HUMAN
US-09-269-250E-38
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-744-549-15
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APPLICANT: Li, Xiao-Jiang
APPLICANT: Li, Shi-Hua
APPLICANT: Li, Shi-Hua
APPLICANT: Sharp, Alan
APPLICANT: Sharp, Alan
APPLICANT: Horley, Paul
APPLICANT: Morley, Paul
APPLICANT: Snyder, Solomon
TITLE OF INVENTION: Huntingtin-associated protein
FILE REFERENCE: 01107.52271
CURRENT APPLICATION UMMBER: US/08/556,419C
CURRENT FILING DATE: 1995-11-09
NUMBER OF SEQ ID NOS: 25
                                                                                                                                                                                                                                                                                                                  US-09-744-549-15
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Best Local Similarity
Thehes 5; Conserve
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                                                                                                      GENERAL INFORMATION:
APPLICANT: Itoh, Kyogo et al.
APPLICANT: Itoh, Kyogo et al.
TITLE OF INVENTION: HLA-A2-restricted Tumor Antigen Peptides Derived From SART-1
FILE REFERENCE: 0020-4808P
CURRENT APPLICATION NUMBER: US/09/744,549
CURRENT FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: JP H10-212940
PRIOR FILING DATE: 1998-07-28
NUMBER OF SEQ ID NOS: 40
COUNTRIES DESCRIPTIONS: 40
RUMBER OF SEQ ID NOS: 40
                                                                        SOFTWARE: PatentIn version 3.0 SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn version 3.1
SEQ ID NO 38
                                                                                                                                                                                                                                                                                 Sequence 15, Application US/09744549 Patent No. 6664232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 4; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 7
TYPE: PRT
                                                       ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 22;
Pred. No.
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Vandekerckhove, Joel
APPLICANT: Gevaert, Kris
TITLE OF INVENTION: METHODS AND APPARATUS FOR GEL-FREE QUALITATIVE AND
TITLE OF INVENTION: QUANTITATIVE PROTEOME ANALYSIS, AND USES THEREFOR
FILE REFERENCE: VBV-001
CURRENT APPLICATION NUMBER: US/10/394,980
CURRENT FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: W50/278,171
PRIOR APPLICATION NUMBER: US60/278,171
PRIOR APPLICATION NUMBER: US60/318,749
PRIOR FILING DATE: 2001-09-12
PRIOR APPLICATION NUMBER: US60/323,999
PRIOR FILING DATE: 2001-09-20
NUMBER OF SEQ ID NOS: 473
                                       TITLE OF INVENTION: METHODS AND APPARATUS FOR GEL-FREE QUALITATIVE AND TITLE OF INVENTION: QUANTITATIVE PROTEOME ANALYSIS, AND USES THEREFOR FILE REFERENCE: VBV-001

CURRENT APPLICATION NUMBER: US/10/394,980

CURRENT FILLING DATE: 2003-03-21

PRIOR APPLICATION NUMBER: PT/PO2/03368

PRIOR FILING DATE: 2002-03-22

PRIOR PRIOR APPLICATION NUMBER: US60/278,171

PRIOR APPLICATION NUMBER: US60/278,171

PRIOR APPLICATION NUMBER: US60/318,749

PRIOR PILING DATE: 2001-09-12

PRIOR PILING DATE: 2001-09-12

PRIOR PILING DATE: 2001-09-20

NUMBER OF SEQ ID NOS: 473
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US-10-394-980-464
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US-10-394-980-421
                                                                                                                                                                                                                                                                                                                                                                       Sequence 464, Application US/10394980 Patent No. 6908740
GENERAL INFORMATION:
APPLICANT: Vandekerckhove, Joel
APPLICANT: Gevaert, Kris
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Best Local Similarity
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SOFTWARE: PatentIn version 3.1
EQ ID NO 464
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: MISC FEATURE
OTHER INFORMATION: FR10 - 1624,68
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ORGANISM: Homo sapiens
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o. 690874
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71.4%;
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71.4%;
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
0; Mismatches 2
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US-08-459-568-20
                                                          RESULT 13
US-08-399-411-20
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                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/399,411
FILING DATE: 06-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1264
TELECOMFUNICATION INFORMATION:
TELEPHONE: (619) 535-9901
TELEPHONE: (619) 535-9901
TELEPAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20, Appli
Patent No. 5811304
Sequence 20, Application US/08399411 Patent No. 5831008 GENERAL INFORMATION:
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                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION: APPLICANT: Huang,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 9
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: MISC FEATURE
OTHER INFORMATION: Table XIX - 1624.68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/459,568
APPLICATION NUMBER: US/08/459,568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  y Match 59.5%;
Local Similarity 71.4%;
les 5; Conservative
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 02-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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                                                                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: linear
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4370 La Jolla Village Drive, Suite 700
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
1; Mismatches 0
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US-08-516-859A-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/399,411
PILING DATE: 06-MAR-1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1264
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20, Application US/08516859A
Patent No. 6069231
GENERAL INFORMATION:
APPLICANT: Huang, Shi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentln Release #1.0, V6
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/516,859A
FILING DATE: 18-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 3
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TITLE OF INVENTION: Retinoblastoma Protein -
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 93
CORRESSEN: Campbell and Flores
ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite
                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
FILING DATE: 06-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0:
                                                   CLASSIFICATION: 514
PRIOR APPLICATION NUMBER: US 08/399,411
                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
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ZIP: 92122
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STATE: California
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STATE: California
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milarity 80.0%;
Conservative
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4370 La Jolla Village Drive, Suite 700
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                                                                                                                                    US/08/516,859A
 US 08/292,683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 21; DB 1;
Pred. No. 4.6e+05;
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; SEQUENCE DESCRIPTION: SEQ ID NO: 20: US-09-586-472-20
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US-09-586-472-20
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ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 1776
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-8949
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 S-09-386-412-20
Sequence 20, Application US/09586472
Sequence 20, Application US/09586472
Patent No. 6323335
GENERAL INFORMATION:
APPLICANT: Huang, Shi
APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
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Best Local Similarity
Matches 4; Conserv
                                                                                                      TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                            APPLICATION NUMBER: US 09/528,706
FILING DATE: 17-MAR-2000
APPLICATION NUMBER: US 08/516,859
FILLING DATE: 18-AUG-1995
APPLICATION NUMBER: US 08/399,411
FILLING DATE: 18-AUG-1995
APPLICATION NUMBER: US 08/292,683
FILING DATE: 18-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 4130
TELEOPHONE: (619) 535-9001
TELEOPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/586,472
PILING DATE: 01-Jun-2000
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
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COUNTRY: USA
                                             TYPE: amino acid
TOPOLOGY: linear
                                                                                    LENGTH: 6 amino acids
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    Mismatches

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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Gapop 10.0 , Gapext 0.5
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1 VLXDDLLEA 9
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Published Applications AA Main:*

1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US11B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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 18-10-861-335-1
18-11-007-740-18
18-11-007-740-29
18-11-007-740-29
18-11-623-176-42
18-10-623-176-45
18-10-623-176-45
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18-10-623-176-45
18-10-623-176-47
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18-10-623-176-3
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Similarity 9; Conservat VLXDDLLEA 9	עם	ANT: GOULMY, ELS A.J.M. ANT: GOULMY, ELS A.J.M. ANT: Hunt, Donald F. ANT: Engelhard, Victor OF INVENTION: HA-1 epit REFERENCE: 2183-6047US IT APPLICATION NUMBER: UP APPLICATION NUMBER: 09/ FILING DATE: 2000-01-21 APPLICATION NUMBER: EP FILING DATE: 1997-07-23 APPLICATION NUMBER: PCT FILING DATE: 1998-07-23 APPLICATION NUMBER: J PRILING DATE: 1998-07-23 APPLICATION NUMBER: J FILING DATE: 2000-01-24 COF SEQ ID NOS: 77 RE: PATENTIN Ver. 2.1	A 1 Application US/10623176	50 50 50 50 50 50 50 50 50 50 50 50 50 5
97.3% 100.0 :ive	l Sequence Descriptio Wherein X	Els A.J.M. Donald F. HA-1 epitop 3-6047US 1 NUMBER: US/S: 2003-07-1 Ver. 2.1	US/10 92446	o o o o o o o o o o o o o o o o o o o
%; Score 36; DB 4; Length 9; 0%; Pred. No. 1.7e+06; 0; Mismatches 0; Indels	.l Sequence Description of Artificial Sequence: wherein X can be R or H	J.M. F. F. F. P. Color H. Colo	ALIGNMENTS 623176 A1	US-10-394-980-464 US-10-275-652-50 US-10-952-557-464 US-10-952-557-464 US-10-96-306-109 US-09-833-203-54 US-10-057-4758-10608 US-10-057-4758-10813 US-10-154-8848-10813 US-10-154-8848-10813 US-10-154-8848-10813 US-10-154-8848-10813 US-10-154-8848-10813 US-10-154-8848-10813 US-10-154-8848-10813 US-10-154-8848-10813 US-10-154-8848-10813 US-10-154-83-176-14 US-10-791-217-4 US-10-809-790-23 US-10-99-364-73 US-11-097-740-40 US-11-07-740-40 US-11-166-225A-168 US-10-166-225A-169 US-10-166-225A-169 US-10-166-225A-169
9; els 0; Gaps	HA-1 peptide	, ,		Sequence 464, Apsequence 421, Apsequence 421, Apsequence 109, Apsequence 10608, Sequence 10603, Sequence 10813, Sequence 10813, Sequence 14, Appsequence 23, Appsequence 23, Appsequence 40, Appsequence 416, Appequence 416, Appsequence 416, Appsequenc
0;				4, App , App 1, App 1, App 9, App 9, App 9, App 9, App 1608, A 813, A 814, A 815, A 81

RESULT 2
US-10-623-176-2
; Sequence 2, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
; APPLICANT: Goulmy, Els A.J.M.
; APPLICANT: Hunt, Donald F.

APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US

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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-10
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Best Local Similarity 85...
Thes 8; Conservative
                                     Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR FILING DATE: 1997-07-23
                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 09/489,760 PRIOR FILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: HA-1 epitopes and uses thereof FILE REFERENCE: 2183-6047US CURRENT APPLICATION NUMBER: US/10/623,176 CURRENT FILING DATE: 2003-07-18
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TYPE: PRT
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OTHER INFORMATION: Description
FEATURE:
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LOCATION: (1)..(9)
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                                                                                                                                                                                                                                                                  TYPE: PRT
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1 VLXDDLLEA 9
                                         Conservative
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88.9%;
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Pred. No. 1.7e+06;
                                                         Score 36; DB 4;
Pred. No. 1.7e+06;
                                                                                                                                                                                                    of Artificial Sequence: HA-1 peptide
                                           Mismatches
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FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOPTWARE: PATENTIN Ver. 2.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (3)...(3)
; OTHER INFORMATION: Xaa is HISTIDINE OR ARGININE RESIDUE
US-10-791-217-1
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                                                                                    , OTHER INFORMATION: obtained from histocompatibility antigen US-10-791-217-2
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Publication No. US20040191268A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/10791217
Publication No. US20040191268A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 2
Best Local Similarity Matches 8; Conserv
                                             Query Match
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APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                             FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE: OTHER INFORMATION: obtained from histocompatibility antigen
                                                                                                                              ORGANISM: Artificial sequence FEATURE:
                                                                                                                                                                         TYPE: PRT
                                                                                                                                                                                               LENGTH: 9
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      Conservative
                          97.3%; Score 36; DB 4; 1
88.9%; Pred. No. 1.7e+06;
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US-10-791-217-5
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Publication No. US20050031612A1
GENERAL INFORMATION:
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Best Local
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LENGTH: 9
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                                                                                       Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Goulmy, Elsa A.J.M.

TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunot TITLE OF INVENTION: of tumors

FILE REFERENCE: 2183-6479US

CURRENT APPLICATION NUMBER: US/10/861,335

CURRENT FILING DATE: 2004-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: obtained from KIAA0223 partial complementary DNA
-10-791-217-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/NL02/00791
PRIOR FILING DATE: 2002-12-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                         LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial sequence
FEATURE:
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8; Conserv
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8
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                                   VLXDDLLEA 9
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VLHDDLLEA
                                                                       Conservative
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Pred. No. 1.7e+06;
0; Mismatches 1
                                                                    Score 36; DB 5;
Pred. No. 1.7e+06;
0; Mismatches 1
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RESULT 8
US-11-007-740-18
J. Sequence 18, Application US/11007740
Publication to US20050233350A1
GENERAL INFORMATION:
APPLICANT: GOLLINY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FOR TREFERENCE: 2799/58994-A
TYPE REFERENCE: 2799/58994-A
TYPE REFERENCE: 2799/58994-A
RESULT 10
US-11-007-740-29
; Sequence 29, Application US/11007740
; Publication No. US20050233350A1
; GENERAL INFORMATION:
; APPLICANT: Goulmy, Elsa
; TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
; FILE REFERENCE: 2799/58994-A
; CURRENT APPLICATION NUMBER: US/11/007,740
; CURRENT FILING DATE: 2004-12-08
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Sequence 20, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
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Best Local S
Matches 8
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TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR PILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin version 3.1
SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 8; Conserv
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CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
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Pred. No. 1.
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Pred. No. 1.7e+06;
0; Mismatches 1
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1.7e+06;
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                                                                                                         US-10-623-176-42
                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 7; Conserva
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LENGTH: 9
TYPE: PRT
             Sequence 42, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 41, Application US/106
Publication No. US20040092446A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor
APPLICANT: Engelhard, Victor H.
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                                                                                                                                                                                                                                                                                                                                                                                                FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                              FEATURE:
NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                               LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENGTH:
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                                                                                                                                                                                 2 VLHDDLLE
                                                                                                                                                                                                                    1 VLXDDLLE 8
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Conservative 0
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                                                                                                                                                                                                                                                                       86.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
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. 1.7e+06;
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hes 0;
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APPLICANT: GOLIMY, Els A.J.M.

APPLICANT: HUNT, Donald F.

APPLICANT: Engelhard, Victor H.

APPLICANT: Engelhard, Victor H.

APPLICANT: Engelhard, Victor H.

APPLICANT: Engelhard, Victor H.

FILE REFERENCE: 2183-6047US

CURRENT APPLICATION NUMBER: US/10/623,176

CURRENT FILING DATE: 2003-07-18

PRIOR APPLICATION NUMBER: 09/489,760

PRIOR APPLICATION NUMBER: BP 97202303.0

PRIOR APPLICATION NUMBER: BP 97202303.0

PRIOR APPLICATION NUMBER: BP 97202303.0

PRIOR APPLICATION NUMBER: DF 707-23

PRIOR APPLICATION NUMBER: DCT/NL98/00424

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR APPLICATION DATE: 1998-07-23

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR APPLICATION DATE: 2000-01-24

NUMBER OF SEQ ID NOS: 77

SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 13
US-10-623-176-45
; Sequence 45, Application US/10623176
; Publication No. US20040092446A1
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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-42
                                                                                                                                                          ; LOCATION: (1)..(9) US-10-623-176-45
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-24
PRIOR FILING DATE: 1998-07-24
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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Best Local
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                                                                       Matches
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                                                                                                                                                                                  FEATURE:
NAME/KEY: SITE
LOCATION: (1)..
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                                                                                                                                                                                                                                                                                                                            LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                               FEATURE:
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Local Similarity 87.5%;
es 7; Conservative
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2 LXDDLLEA 9
                                                                       Similarity 7; Conserv
                                                                         Conservative
                                                                                            86.5%;
87.5%;
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Pred. No. 1.7e+06
                                                                                                 Score 32; I
Pred. No. 1
                                                                            Mismatches
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US-10-623-176-47

Sequence 47, Application US/10623176

Publication No. US20040092446A1

GENERAL INFORMATION:
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR PILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
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US-10-623-176-46
US-10-623-176-46
; Sequence 46, Application US/10623176
; Publication No. US20040092446A1
; GENERAL INFORMATION:
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
                                                                                                                                    NUMBER OF SEQ ID NOS: 77
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 47
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Best Local Similarity
Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -10-623-176-46
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OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
FEATURE:
NAME/KEY: SITE
LOCATION: (1)..(9)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                        TYPE: PRT
                                                                                                         LENGTH:
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Search completed: April 6, 2006, 16:50:29 Job time: 91.3333 secs THIS PAGE BLANK (USPTO)

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                               Database :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence:
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Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/USOS NEW PUB.pep:*

2: /SIDSS/ptodata/2/pubpaa/USOS NEW PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/USO7 NEW PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT NEW PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/USO9 NEW PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*

7: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*

8: /SIDSS/ptodata/2/pubpaa/USO1 NEW PUB.pep:*
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37
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Gapop 10.0 , Gapext 0.5
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	B	ID	Description
1	36	97.3	9	7	US-11-010-748A-11	Sequence 11, Appl
N	36	97.3	9	7	-11-010-748A-	
w	21	56.8	9	0	-10	63
4.	21	56.8	9	σ	US-10-927-634-63	63,
5	20	54.1	v	7	US-11-129-143-168	168,
0	20	54.1	ហ	7	US-11-129-143-169	169,
7	20	54.1	₅	7	US-11-129-143-170	170,
8	20	54.1	ហ	7	US-11-129-143-171	171,
9	20	54.1	₅	7	US-11-129-143-172	
10	20	54.1	v	7	US-11-129-143-173	173,
11	20	54.1	7	σ	US-10-982-440-126	126,
12	20	54.1	8	7	US-11-021-305-62	62,
13	20	54.1	80	7	US-11-021-305-65	65,
14	20	54.1	80	7	-305	Sequence 66, Appl
15	19	51.4	Φ	7	US-11-207-078-373	373,
16	19	51.4	7	7	US-11-225-686-2954	
17	19	51.4	7	7	US-11-225-686-3877	3877,
18	19	51.4	7	7	US-11-225-686-3914	3914,
19	19	51.4	7	7	US-11-225-686-3915	
20	19	51.4	7	7	US-11-225-686-3916	3916,
21	19	51.4	7	7	US-11-225-686-3932	
22	19	51.4	7	7	US-11-225-686-3941	3941,
23	19	51.4	7	7	-009	2954,
24	19	51.4	7	7	-009	
25	19	51.4	7	7	170 11 303 000 3014	301

RESULT 2 US-11-010-748A-12 Sequence 12, Ap Publication No. GENERAL INFORMA APPLICANT: MET APPLICANT: MOT APPLICANT: MOT APPLICANT: SC APPLICANT: SC	Query Matc Best Local Matches Qy	PRIOR FILL PRIOR PILL PRIOR FILL PRIOR FILL UNMBER OF SOFTWARE: SEQ ID NO 1 LENGTH: 9 I LENGTH: 9 I TYPE: PRI ORGANISM: FEATURE: FEATURE: FEATURE: OTHER INFO	ESULT 1 S-11-011 Sequence Publica GENERAL APPLIC APPLIC APPLIC APPLIC TITLE FILE F CURREN CURREN		2222 2222 3222 322 322 322 322 322 322
748A-12 12, Applicat 10n No. US2 10	cch al Similarity al Similarity b; Conservat 1 VLXDDLLEA 9 1 VLHDDLLEA 9	ING DATE: 3 ING DATE: 3 SEQ ID NOS SEQ ID NOS PatentIn v 1 Homo sapi Homo sapi ORMATION: (1) . (2) ORMATION: (3) . (4) . (5)	SULT 1 -11-10-748A-11 -11-10-748A-11 Sequence 11, Application US/ Publication No. US2005024442 GENERAL INFORMATION: APPLICANT: Merck Patent Gmb APPLICANT: MERCK PATENT Gmb APPLICANT: MOLL, Heidrun APPLICANT: SCHARM, Burkhar TITLE OF INVENTION: COMPOSI FILE REFERENCE: MER-136 CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 2004-		19 51.4 19 51.4 18 48.6 18 48.6
ion US/1101074 50244421A1 ent GmbH TTTER, Wolfgang	97.3%; E 88.9%; E ive 0;	13 13 13 3.1 3.1	11010 1A1 H olfgal olfgal d d US/1:		7 7 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
3 18A	icore 36; DB 7; red. No. 1.4e+05 Mismatches	13423.5 histocompatibility antigens	HODS 8A	ALIGNMENTS	US-11-202-009-3915 US-11-202-009-3932 US-11-202-009-3932 US-11-202-009-3932 US-11-202-009-3931 US-10-927-03-39 US-10-989-767A-39 US-10-927-634-62 US-11-96-706-88 US-11-96-706-88 US-11-96-706-89 US-11-98-767A-219 US-10-989-767A-219 US-10-989-767A-219 US-10-989-767A-3163 US-11-129-741-363 US-11-129-741-363 US-11-129-143-163
	Length 9; ; 1; Indels		FOR MODULATING IMMUNE		
	0; Gaps	HA-1 characterized	MMUNE RESPONSE		e 3915, e 3932, e 3932, e 3941, pe 7, App e 62, App e 62, App e 62, App e 62, App e 62, App e 1363, P e 1363, P e 1363, P e 1363, P
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RESULT 3
US-10-927-435-63
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Publication No. US20060045884A1
GENERAL INFORMATION:
APPLICANT: MOLLDREM, JEFFREY
TITLE OF INVENTION: VACCINES FOR AUTOIMMUNE AND INFECTIOUS DISEASE
FILE REFERENCE: UTSC:788USC2
CURRENT FILING DATE: 2004-08-26
PRIOR APPLICATION NUMBER: US/10/927,435
CURRENT FILING DATE: 2004-08-26
PRIOR APPLICATION NUMBER: 60/489,238
PRIOR FILING DATE: 2003-08-26
NUMBER OF SEQ ID NOS: 63
                     Sequence 63, Application US/10927634
Publication No. US20060045881A1
GENERAL INFORMATION:
APPLICANT: MOLLDREM, JEFFREY
TITLE OF INVENTION: ANTI-CANCER VACCINES
FILE REFERENCE: UTSC:788USC1
CURRENT APPLICATION NUMBER: US/10/927,634
CURRENT FILING DATE: 2004-08-26
CURRENT FILING DATE: 2004-08-26
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PRIOR FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: EP02013423.5
PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
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CURRENT FILING DATE: 2004-12-13
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TYPE: PRT
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NAME/KEY: MISC FEATURE
LOCATION: (1)...(9)
OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by T
OTHER INFORMATION: ell epitopes
PRIOR APPLICATION NUMBER: 60/489,238
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
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88.9%;
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Pred. No.
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Pred. No. 1.4e+05;
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APPLICANT: YELLSEBY, ANNE F.

TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 169
LENGTH: C.
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US-11-129-143-168
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APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Wei
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APPLICANT: BRETZEL, Wer
APPLICANT: HUMBELIN, Ma
APPLICANT: LOPEZ-ULIBAN
APPLICANT: MAYER, Anne
APPLICANT: YELISEEV, Al
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SEQ ID NO 168
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Best Local Similarity
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Matches 5; Conserv
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TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT PEPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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TYPE: PRT
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                             BRETZEL, Werner
HUMBELIN, Markus
LOPEZ-ULIBARRI, Rual
MAYER, Anne F.
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LOPEZ-ULIBARRI, I
MAYER, Anne F.
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Pred. No. 1.4e
1; Mismatches
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Pred. No. 1.4e+05;
2; Mismatches C
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APPLICANT: HUMBELIN, Markus
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: YELISEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C198435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: Patentin version 3.1
SEQ ID NO 171
LENGTH: 5
TYPE: PRT
ORGANISM: Bacillus subtilis
US-11-129-143-171
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US-11-129-143-170
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US-11-129-143-169
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Best Local Similarity 60.0
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                                                                             Query Match
Best Local &
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                                                                 Matches
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APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
RUMBER OF SEQ ID NOS: 197
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APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
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             4 DDLLE 8
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Pred. No.
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Pred. No. 1.4e+05;
2; Mismatches 0
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                                                                             1.4e+05;
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APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: PatentIn version 3.1
SEQ ID NO 172
LENGTH: 5
CRGANISM: Bscherichia coli
RESULT 11
US-10-982-440-126
(S-10-982-440-126); Sequence 126, Application US/10982440
; Publication No. US20060018909A1
; GENERAL INFORMATION:
; APPLICANT: Oliner, John
                                                                                                                                                                                          Query Match
Best Local Similarity
Watches 3; Conserve
                                                                                                                                                                                                                                                                      ; TYPE: PRT
; ORGANISM: Haemophilus influenzae
US-11-129-143-173
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US-11-129-143-172
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; Publication No. US200
; GENERAL INFORMATION:
; APPLICANT: BERRY, Al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 60.0
                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: BERRY, Alan
APPLICANT: BERRY, SAlan
APPLICANT: HUMBELIN, Markus
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.1 SEQ ID NO 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 173, Application US/11129143 Publication No. US20050266518A1
                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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                                                                                                                                                                                                                                                                                                                            ENGTH: 5
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                                                                                                                                                                                                             Conservative
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Pred. No. 1.4e+05;
2; Mismatches 0;
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Pred. No. 1.4e+05;
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RESULT 13
US-11-021-305-65
; Sequence 65, Application US/11021305
; Publication No. US20050282733A1
; GENERAL INFORMATION:
    APPLICANT: Hutley, Louise J
; TITLE OF INVENTION: Differentiation-modulating agents and uses therefor FILE REFERENCE: DAVI169.001CP1
; CURRENT APPLICATION NUMBER: US/11/021,305
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: USSN 60/392,130
; PRIOR FILING DATE: 2002-06-27
; PRIOR FILING DATE: 2002-06-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 12
US-11-021-305-62
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CURRENT FILING DATE: 2004-11-04
PRIOR APPLICATION NUMBER: 60/620,161
PRIOR FILING DATE: 2004-10-19
NUMBER OF SEQ ID NOS: 215
SOFTWARE: Patentin version 3.3
SEQ ID NO 126
LENGTH: 7
LENGTH: 7
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Publication No. US20050282733A1
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Matches 4; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn version 3.3 SEQ ID NO 62
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                                                                    NUMBER OF SEQ ID NOS: 170
SOFTWARE: PatentIn version 3.3
SEQ ID NO 65
LENGTH: 8
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APPLICANT: Hutley, Louise J
TITLE OF INVENTION: Differentiation-modulating agents and uses therefor
FILE REFERENCE: DAVI169.001CP1
CURRENT APPLICATION NUMBER: US/11/021,305
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR FILING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 170
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TITLE OF INVENTION: Angiopoietin-2 Specific Binding Agents
FILE REFERENCE: 04-881-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 8
TYPE: PRT
ORGANISM: mammalian
ORGANISM: mammalian
                                                     TYPE: PRT
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100.0%; Pred. No. 1.4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         54.1%; Score 20; DB 7; L
100.0%; Pred. No. 1.4e+05;
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CURRENT APPLICATION NUMBER: US/11/021,305
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: USSN 60/392,130
PRIOR FILING DATE: 2002-06-27
NUMBER OF SEQ ID NOS: 170
SOFTWARE: Patentin version 3.3
SEQ ID NO 66
LENGTH: 8
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US-11-207-078-373
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US-11-021-305-66
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Best Local Similarity
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Publication No. US20060040307A1
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APPLICANT: Hutley, Louise J
TITLE OF INVENTION: Differentiation-modulating agents and uses therefor FILE REFERENCE: DAVII69.001CP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Cech
COMPUTER READABLE FORN:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/11/207,078
FILING DATE: 17-Aug-2005
CLASSIFICATION LOTA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION UMBER: US/09/721,477
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Local Similarity 100.0%; Pred. No. 1.4e+05;
                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                          Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 DDLL 7
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STATE: California
                                                                                                                                                                                                                                              COUNTRY: USA
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Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
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100.0%; Pred. No. 1.4e+05;
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PELLIANTION NUMBER: 03/944,549

PELLIAN DATE: dhinown.

PELLIAN DATE: 18-APR.197

PELLIAN DATE: 25-APR.197

PAPLICATION NUMBER: US 06/844,017

PELLIAN DATE: 25-APR.197

PAPLICATION NUMBER: US 06/854,050

PELLIAN DATE: 06-MAY.197

PAPLICATION NUMBER: US 06/951,93

PELLIAN DATE: 14-AUG.197

PAPLICATION NUMBER: US 06/951,503

PELLIAN DATE: 01-OCT.1997

APPLICATION NUMBER: US 06/951,503

PELLIAN DATE: 01-OCT.1997

PAPLICATION NUMBER: 01-OC
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Minimum DB
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                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: geneseqp1980
2: geneseqp2000
3: geneseqp2001
4: geneseqp2001
5: geneseqp2003
7: geneseqp2003
8: geneseqp2004
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Query
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Match 100%
first 45 summaries
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ADS78893	ADT39474	AAE28095	AA021066	AAB70589	AAW33386	AAR69140	AAR29358	ADC25889	AAE26631	AAB84492	AAG79144	ABU97230	ABU97229	ADU99636	ADT49074	ADT73023	ADM96258	ADM96261	AAB31275	ADM96262
Ads78893 SARS viru	Adt39474 hSARS vir	Aae28095 Human imm	Aao21066 Isomerise	Aab70589 Human imm	Aaw33386 Altered I	Aar69140 Endotheli	Aar29358 Endotheli	Adc25889 Yeast GPA	Aae26631 Yeast GPA	Aab84492 Peptide f	Aag79144 Amino ter	Abu97230 Enzyme pe	Abu97229 Enzyme pe	Adu99636 BFA5 tumo	Adt49074 Human BFA	Adt73023 Human RSV	Adm96258 Human ser	Adm96261 Human ser	Aae31275 Human mag	Adm96262 Human ser

AAW99196 standard; peptide; 9 AA.

ALIGNMENTS

RESULT 1
AAW99156
ID AAW9
XX AAW9
XX Minc
XX Homc
XX Homc
XX Hom
X AAW99196; 20-MAY-1999 (first entry)

Minor histocompatibility antigen HA-1 T-cell epitope #2

Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.

Homo sapiens.

04-FEB-1999. WO9905174-A1.

23-JUL-1998; 98WO-NL000425.

23-JUL-1997; 97EP-00202303

(UYLE-) RIJKSUNIV LEIDEN

Goulmy EAJM, Hunt DF, Engelhard VH;

WPI; 1999-153312/13.

A new minor histocompatibility antigen, HA-1 - useful to treat immune diseases and prevent rejection and host versus graft disease in bone marrow and organ transplantation.

Claim 3; Page 32; 47pp; English.

The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the diseases reatment of severe aplastic anaemia, leukaemia, and immune deficiency

Sequence 9 AA;

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RESULT 3
ADH40333
ID ADH4
XX
AC ADH4
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                         ADH40333 standard; peptide; 9 AA.
                                                                                                                                                                                                      The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
ADH40333;
                                                                                                                                                                                  Sequence 9
                                                                                                                                                                                                                                                                                                                                            Claim 18; Fig 5;
                                                                                                                                                                                                                                                                                                                                                                    of genetic aberrances.
                                                                                                                                                                                                                                                                                                                                                                                 Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. de
                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-142960/12
                                                                                                                                                                                                                                                                                                                                                                                                                                            Goulmy
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02-JUN-1998;
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98EP-00870125.
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                                                                                                                                                                                                                                                                                                                                            59pp; English.
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Pred. No.
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Pred. No. 2e+06;
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2e+06;
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RESULT 4
AAW99197
ID AAW9
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Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
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                                                                                                 Minor histocompatibility antigen HA-1 T-cell epitope #3
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Pred. No. 2e+
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RESULT 5
AAW97374
ID AAW9
XX AAW
DT 13-1
DT 13-1
XX Int
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    WPI; 1999-142960/12
                                                Goulmy E;
                                                                                                                                              23-JUL-1997;
02-JUN-1998;
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                                                                                               (UYLE-) RIJKSUNIV LEIDEN
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88.9%;
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Pred. No.
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2e+06;
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versus
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RESULT 6
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ID ADH4
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The invention relates to a novel method for providing epitopes of allelic variants of antigenic proteins from specific species based on single nucleotide polymorphism (SNP), by defining target protein/peptide or its subset, screening database of DNA encoding target protein, identifying, selecting allelic peptide/protein variants, expression product or its fragment encoded by DNA sequence having SNP, creating variant epitopes, selecting epitopes binding to MHC protein. A protein of the invention has cytostatic activity, and may have a use in a vaccine. The method is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the CDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplatic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                          Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                 Strittmatter W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human; cytostatic; vaccine; SNP profile; minor histocomptibility antigen; mHAg; T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Typing minor histocompatibility antigen HA-1 - identifying HA-1 R and HA-1 H alleles of HA-1, of genetic aberrances.
                                                                                                                                                          Disclosure; Page 82; 119pp; English.
                                                                                                                                                                                                                                                                                 WPI; 2004-082200/08
                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUN-2003; 2003WO-EP006251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human minor histocomptibility antigen HA-1 T cell eptiope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH40334;
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                                                                                                                                                                                                                                                                                                                                                                                  13-JUN-2002; 2002EP-00013423
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                                                                                                                                                                                                                                                                                                                                                  (MERE ) MERCK PATENT GMBH.
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                                                                                                                                                                                                                                                                                                                 Moll H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vaccine; SNP profile; cancer; leukaemia;
ility antigen; mHAg; T cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peptide;
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Pred. No.
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2e+06;
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                           The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. Pl is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                                                                                                         A new minor histocompatibility antigen, dispasses and prevent rejection and host
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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Sequence 9 AA
                                                                                                                                                                                                                                                                                                       Goulmy EAJM, Hunt DF,
                                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-1999
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                                                                                                                                                                         Claim 1; Page 32; 47pp; English
                                                                                                                                                                                                                                                                                                                                                                        23-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                    23-ЛИТ-1998;
                                              treatment of severe aplastic anaemia, leukaemia, and immune
                                                                                                                                                                                                                                                                                                                                       (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                           1999-153312/13.
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                                                                                                                                                                                                          and organ transplantation.
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8; Conserv
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Pred. No. 2e+0
0; Mismatches
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2e+06;
                                                                                                                                                                                                                       HA-1 - useful to treat immune versus graft disease in bone
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                                                in the deficiency
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RESULT 8
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ID AAW9
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Best Local S
Matches 8
                                                                       Query Match
                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune dis neoplastic haematopoietic cell.
                                                                                                                                The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen HA-1. The peptide can be used in vaccines or pharmaceutical formulations as medicines to induce tolerance for transplants so as to prevent rejection and/or Graff-versus-Host Disease, or to treat autoimmune diseases.

Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides. The peptides can also be used to raise antibodies, T-cell receptor, B-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T-cell epitope from the minor histocompatibility antigen
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                                                                                                Sequence 9 AA;
                                                                                                                                                                                                                                                                     Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-versus-host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAY-1999 (first entry)
                                                                                                                         The peptides can also be used and T-cells
                                                                                                                                                                                                                                               Claim 1; Page 39; 57pp; English
                                                                                                                                                                                                                                                                                                                        WPI; 1999-142855/12.
                                                                                                                                                                                                                                                                                                                                                                                               23-JUL-1997;
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                                                           Local
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                                                                                                                                                                                                                                                                                                                                               EAJM,
                        1 VLHDDLLEA 9
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8; Conser
                                                8;
                                                             Similarity
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  VLXDDLLEA
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                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                               DF,
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Pred. No. 2e+06;
                                                              Score 36;
Pred. No.
                                                  Mismatches
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                                                              DB 2;
2e+06;
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RESULT 9 ADU99853 ID ADU9

ADU99853 standard; peptide; 9 AA.

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ARBSULT 10
AAW97373
ID AAW97
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                                                                                       Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease.
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          Misc-difference
                                 Misc-difference
                                                                                                                                      Peptide
                                                                                                                                                             13-MAY-1999
                                                                                                                                                                                  AAW97373;
                                                                                                                                                                                                     AAW97373 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                          Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                 the
                                                                                                                                                                                                                                                                                                                                                                                                                            encoding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berinstein N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-MAY-2003;
16-MAY-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BCZ4 tumor antigen antigenic peptide #85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-FEB-2005
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                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                               invention comprises an expression vector that contains a nucleic acid oding a tumor antigen (e.g. BCZ4 or BFY3). The expression vector of invention is useful for the expression of multiple tumor antigens and prevention or treatment of cancer. The present amino acid sequence resents a BCZ4 tumor antigen antigenic peptide that was used in an mple of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expression vector comprising a nucleic acid encoding a tumor antigen, . BFA4, BCY1, BFA5, BCZ4, or BFY3, useful for expressing multiple or antigens, or for preventing or treating cancer.
                                                                  sapiens
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                                                                                                                                     epitope
                                                                                                                                                                                                                                                                                                                                                                                                                                                            8; Page 60; 109pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AVENTIS PASTEUR INC
                                                                                                                                                                                                                                                                 LHSDLLE 8
                                                                                                                                                                                                                                                                                       THODILE 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide.
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                                                                                                                                                                                                                                                                                                             Conservative
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2003US-0471193P.
                                                                                                                                                           (first entry)
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                                   Location/Qualifiers
                                                                                                                                     of HA-1 antigen.
/note= "not specified"
                      /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumor antigen; cancer; cytostatic; BCZ4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                        68.9%;
85.7%;
                     "not
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
                     specified"
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Pred. No. 2e+0
0; Mismatches
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2e+06;
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Best Local S
Matches 6
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                                                                                                                                                    Cytotoxic T-lymphocyte response; CTL; antigen; lymphatic immunisation; tumour; infectious disease; immunotherapy; malignant melanoma; viral disease; hepatitis; AIDS.
           10-JUL-1997;
10-DEC-1997;
                                                                                                                              Synthetic.
                                                                                                                                                                                                      T cell
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                                             10-JUL-1998;
                                                                     21-JAN-1999
                                                                                            WO9902183-A2
                                                                                                                                                                                                                                                      AAY10122;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of genetic aberrances.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Typing minor histocompatibility antigen HA-1 - by amplifying identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g.
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02-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Goulmy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9905313-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYLE-) RIJKSUNIV LEIDEN
                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                   herpesvirus
                                                                                                                                                                                                   epitope/MHC
                                                                                                                                                                                                                                                                                                                                                              1 VLHDDLLEA 9
                                                                                                                                                                                                                                                                                                                                                                                      6
                                                                                                                                                                                                                                                                                                                                                                                                                                      ĀΆ;
                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                              (first entry)
           97CA-02209815
97US-00988320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97EP-00202303.
98EP-00870125.
                                             98WO-US014289
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                                                                                                                                                                                                     ligand
                                                                                                                                                                                                                                                                                                                                                                                                 66.7%;
                                                                                                                                                                                                     SEQ
                                                                                                                                                                                                                                                                                                                                                                                   Score 30; DB:
Pred. No. 2e+0
0; Mismatches
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                                                                                                                                                                                                      NO:52
                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                              Length
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Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 method comprises: (a) delivering an antigen to the mammal at a level to induce an immunological CTL response in the mammal; and (b) maintaining the level of the antigen in the mammal; slymphatic system to maintain the immunologic CTL response. The method can be used for the delivery of e.g. a differentiation antigen, a tumour-specific multilineage antigen, an embryonic antigen, an oncogene antigen, a mutated tumour-suppressor gene antigen, or a viral antigen. They can be used for the treatment of disease such as cancer, e.g. mallgnant melanoma or infectious disease, e.g. viral disease such as hepatitis or AIDS. Sustained antigen delivery to the lymphatic system provides for potent CTL stimulation that takes place in the milieu of the lymphoid organ, and it sustains stimulation that the mody. AAY10071 to AAY16339 represent examples of peptide antigens
                                                                                                                                                                                                                                                                                                                                                                06-AUG-2003
15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 25; 199pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inducing a cytotoxic T lymphocyte response - by maintaining a level antigen in the lymphatic system of a mammal so as to provide a sustr CTL response, used to treat, e.g. AIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kuendig TM,
             WPI; 2002-657506/70
                                                                                                                                                                                                                  Cercopithecine
                                                                                                                                                                                                                                                                        antigen;
                                                                                                                                                                                                                                                                                                                                   MHC class I molecule, viral epitope #53.
                                                                                                                                                                                                                                                                                                                                                                                                           ABG79805
                                                                                                                                                                                                                                                                                                                                                                                                                                      ABG79805 standard; peptide; 9 AA.
                                                                                                   02-FEB-2001; 2001US-00776232
                                                                                                                                                                                                                                                                                                      Major histocompatibility complex; MHC; MHC class I molecule; virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunological cytotoxic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CTLI-) CTL
                                                                                                                               22-JAN-2002; 2002WO-US002033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
les 5; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            present invention
                                          MT,
                                                                      CIL
                                                                                                                                                                                                                                                           cytotoxic T lymphocyte response; CTL response; lymphatic system;
immunogenic; malignant tumour; carcinoma; melanoma; leukaemia;
; infectious disease; hepatitis; malaria; measles; tuberculosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ALTHIDDT 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the present invention
                                                                                                                                                                                                                                               immune deficiency
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IMMUNOTHERAPIES CORP
                                          Simard JJL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                      IMMUNOTHERAPIES CORP
                                                                                                                                                                                                                                                                                                                                                                (revised)
(first en
                                                                                                                                                                                                                  herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9
                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  62.2%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            describes a method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T lymphocyte (CTL) response
                                                                                                                                                                                                                                               syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28;
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of inducing and/or sustaining an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in a mammal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a sustained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 13
ADK68732
ID ADK68
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Best Local S
Matches 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is particularly useful for treating a mammal having a malignant tumour (e.g. carcinoma, melanoma, leukaemia or lymphoma) or infectious disease (e.g. hepatitis, acquired immune deficiency syndrome (AIDS), malaria, measles or tuberculosis), or in an animal having a predisposition to these diseases. The mammal may be dogs, cats, mice, cattle, sheep, pigs, goats, rabbits, or preferably humans. ABG79753-ABG80319 represent viral epitopes on major histocompatibility complex (MHC) class I molecules, used in the method of the invention. (Updated 06-AUG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method of inducing and/or sustaining an immunological cytotoxic T lymphocyte (CTL) response in a mammal comprising administering directly to the lymphatic system of the mammal:
(a) an antigen in the form of a polypeptide; (b) a vector comprising a nucleic acid encoding the antigen; or (c) a non-peptide antigen. The method is useful for inducing and/or sustaining CTL response in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           epitope liberation; substrate; proteasome; cytostatic; antibacterial; protozoacide; fungicide; T-cell activator; vaccine; housekeeping epitope; cytotoxic T lymphocyte; CTL; adoptive immunotherapy; neoplastic cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Inducing or sustaining immunological cytotoxic T lymphocyte response i mammal, useful for treating a mammal with malignant tumor or infection disease, by directly administering an antigen to the lymphatic system
                                                                                                                                                                                                                                                                                        (SIMA/)
(DIAM/)
(QIUZ/)
(LEIX/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 19; 73pp;
                                                                                                                                                                                                                               Simard JJL, Diamond
                                                                                                                                                                                                                                                                                                                                                                                                                                          07-NOV-2001; 2001US-0336968P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-NOV-2002; 2002US-00292413.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human herpesvirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Epitope liberation-related peptide SeqID95.
                                                                                                                                                                            WPI; 2004-167209/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2003228634-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  bacterium;
                                                                                                                                                                                                                                                                                                                                               SIMARD J J L.
DIAMOND D C.
                                                                                                                                                                                                                                                                                        QIU Z.
LEI X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protozoan; fungus; housekeeping proteasome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62.2%;
83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    얁
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Disclosure; SEQ ID NO 95; 67pp; English

Identifying polypeptide suitable for epitope e.g., housekeeping epitope. liberation by contacting substrate polypeptide comprising epitope of interest, with proteasome, and assaying for liberation of epitope.

interest, with proteasome, and assaying for liberation

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RESULT 14
ADKOS291
ID ADKOS
XX ADKOS
XX ADKOS
XX PATH
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XX WO1;
XX WATI
PT VIEW
PT 
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Best Local (
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                                                                                                            preventing
                                                                                                                                                                                                                                                                                                                                        Mattner F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-JUL-2002;
11-JUL-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-JUL-2003; 2003WO-EP008112.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HLA-allele; CTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pathogenic virus; alternative reading frame; antigenic determinant; virucide; vaccine; therapeutic agent; infection; epitope peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus CTL epitope peptide #3121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-MAY-2004
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                                                                                                        polypeptide encoded by an alternative reading us comprising an antigenic determinant, useful venting an infection with the pathogenic virus.
                                                                                                                                                                                                                                                               2004-169243/16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VILHDDL 6
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                                                                                                                                                                                                                                                                                                                                        Schmidt W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002AT-00001124.
2003EP-00450171.
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Pred.
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                                                                                                                                               frame of a pathogenic for treating or
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Claim 14; Page 114; 220pp; English

62

Score 28;

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8,

Length

9;

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RESULT 15
ADK05293
ID ADK05
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Best Local S
Matches 5
                                          This invention relates to a novel polypeptide encoded by an alternative reading frame of a pathogenic virus, where the polypeptide starts with a methionine amino acid residue, which comprises an antigenic determinant and more than 7 amino acid residues. The invention may be useful for the production of compounds with a virucide activity or the development of a vaccine. The polypeptide or its fragments may be useful as a therapeutic agent. It is also useful for the manufacture of a medicament for treating or preventing an infection with the pathogenic virus. The present sequence is that of a hepatitis C virus CTL epitope peptide of the
Sequence 9
                                                                                                                                                                                                                                          New polypeptide encoded by an alternative reading frame of a parisus comprising an antigenic determinant, useful for treating preventing an infection with the pathogenic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pathogenic virus; alternative reading
virucide; vaccine; therapeutic agent;
HLA-allele; CTL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention relates to a novel polypeptide encoded by an alternative reading frame of a pathogenic virus, where the polypeptide starts with a methionine amino acid residue, which comprises an antigenic determinant and more than 7 amino acid residues. The invention may be useful for the production of compounds with a virucide activity or the development of a vaccine. The polypeptide or its fragments may be useful as a therapeutic agent. It is also useful for the manufacture of a medicament for treating
                                                                                                                                                                                                             Claim 14; Page 114; 220pp; English.
                                                                                                                                                                                                                                                                                                          WPI; 2004-169243/16.
                                                                                                                                                                                                                                                                                                                                            Mattner F,
                                                                                                                                                                                                                                                                                                                                                                                                           24-JUL-2002; 2002AT-00001124.
11-JUL-2003; 2003EP-00450171.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-JUL-2003; 2003WO-EP008112.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004011650-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hepatitis C virus CTL epitope peptide #3123.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADK05293 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or preventing an infection with the pathogenic virus. The present sequence is that of a hepatitis C virus CTL epitope peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9 AA;
                                 invention.
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0; Mismatches
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infection; epitope peptide;
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2e+06;
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Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 3 HDLL 7

Db 4 HDUL 8

Search completed: April 6, 2006, 16:13:13

Job time: 114.667 secs 144.667 s

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Result
No.
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Maximum DB
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Maximum Match
Listing first
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                Score
                                                                                                                                                                                                                                                                                                                                                                                                                               seq length: 0 seq length: 9
PIR_80:*
1: pir1:*
2: pir2:*
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Match Length
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45
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VLHDDLLEA 9
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          GenCore version 5.1.7 (c) 1993 - 2006 Biocceleration Ltd
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                                                                                                                                                                                                                                                                                        SUMMARIES
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trypsin (EC 3.4.21
T-cell receptor be
biotin A - Citroba
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cerebellar degener
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hucolin, 75K chain
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Έ.	transaldolase (EC	Ig heavy chain CRD	tetrameric protein	ferredoxin a2 - Ja	158K exoantigen -	chitin-binding pro	Ig heavy chain CRD	pilE protein - Esc	phosphoglycerate t	hypothetical prote	copper resistance	T-cell receptor be	T-cell receptor be	late G1-69 protein	pyrimidine synthes

LIGNMENTS

R;Friedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H. Proc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1994
A;Title: Characterization of genomic Clones specifying A;Reference number: 146868; MUID:84221901; PMID:6328491

A;Accession: I46868

alpha-myosin heavy chain - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997

#text_change 09-Jul-2004

rabbit alpha- and beta-ventricul

C;Accession: I46868

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RESULT
PQ0663
                                                                                                                                                                                                                                                                                            phosphoenolpyruvate carboxykinase - Trypanosoma brucei (;Species: Trypanosoma brucei C;Species: Trypanosoma brucei C;Date: 28-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 09-Jul-2004 C;Accession: S55696 R;Hunt, M.; Koehler, F. Biochim. Biophys. Acta 1249, 15-22, 1995 Biochim. Biophys. Acta 1249, 15-22, 1995 A;Title: Purification and characterization of phosphoenolpyruvate carboxykinase from Tr. A;Reference number: S55696; MUID:95284106; PMID:7766679
                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 2
S55696
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A;Molecule type: protein
A;Residues: 1-9 <HUN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-7 <FRI>
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                                                                                                                                         Matches
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                                                                                                   1 АТНООГ 6
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                                                                   IIHKNL 8
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Pred. No. 2.8e+05;
                                                                                                                                                        Score 17; I
Pred. No. 2.
                                                                                                                                         Mismatches
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membrane protein - porcine epidemic diarrhea virus (isolate Belgian CV777) (fragment) C;Species: porcine epidemic diarrhea virus (C;Date: 14-Uul-1994 #sequence_revision 14-Jul-1994 #text_change 08-Oct-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         issible gastroenteritis virus.
A;Reference number: JQ2191; MUID:93389433;
A;Accession: PQ0663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: UNIPARC:UPI0000170FAC; GB:Z14976; C;Comment: This virus is coronavirus related to human C;Keywords: membrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
A; Residues: 1-7 <BRI>
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                                                                                                                                                                           hypothetical protein 8 [imported] - Pseudomonas aeruginosa (fragment) C;Species: Pseudomonas aeruginosa C;Species: Pseudomonas aeruginosa C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 31-Dec-2004 C;Accession: Pc4131 R;Kawasaki, S; Arai, H.; Igarashi, Y.; Kodama, T. R;Kawasaki, S; Arai, H.; Igarashi, Y.; Kodama, T. Gene 167, 87-91, 1995 A;Title: Sequencing and characterization of the downstream region of the ger y for biosynthesis of heme d1. A;Reference number: JC4552; MUID:96144254; PMID:8566817 A;Accession: Pc4131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Homo sapiens (man)
C;Date: 15-Feb-1997 #sequence_revision
C;Accession: S68004
R;Edgar, P.F.
                                                                                                                                                                                                                                                                                                                                                           RESULT 5
PC4131
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: protein A;Residues: 1-7 <EDG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hucolin, 75K chain - human (fragment)
C;Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                                                                                                                                                                   멍
                                                                       A,Cross-references: UNIPROT:P95412; UNIPARC:UPI000017A96A; A,Note: this ORF is not annotated in GenBank entry PSENIRC, C;Superfamily: Pseudomonas stutzeri nirD protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 δ
                                                                                                                               A; Molecule type: DNA
A; Residues: 1-8 < KAW>
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Best Local S
Matches 3
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Best Local Similarity 100.0%; 1

Matches 3; Conservative 0;
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Pred. No.
         Score 16; DB; Pred. No. 2.8
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3. 2.8e+05;
0;
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J. 2.8e+05;
O;
     DB 2;
2.8e+05;
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                                            Length 8;
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S78764
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A59489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Cross-references: UNIPAL
C; Keywords: mitochondrion
F;1-6/Product: ribosomal 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     submitted to the Protein Sequence
A;Reference number: S78760
A;Accession: S78764
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C;Species: Rattus norvegicus
C;Date: 25-Aug-2003 #sequence_revision 25-Aug-2003 #text_change 25-Aug-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: prot
A; Residues: 1-6 < GRA>
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S20446
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                                                                                                   R;Kessler, E.; Safrin, M.; Peretz, M.; Burstein, Y. FEBS Lett. 299, 291-293, 1992
A;Title: Identification of cleavage sites involved in A;Reference number: S20446; MUID:92183956; PMID:154450
A;Accession: S20446
                                                                                                                                                                                                              elastase - Pseudomonas aeruginosa
C;Species: Pseudomonas aeruginosa
C;Date: 19-Mar_1997 #sequence_revision 24-Jul-1997 #text_change
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A; Residues: 1-7 < NEG>
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A;Accession: A59489
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                                                   A;Molecule type: protein
A;Residues: 1-7 <KES>
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                                  A; Cross-references:
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                                      UNIPARC: UPI000017A95F
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     31.1%; Score 14;
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venom protein HR-3 - oriental hornet (fragment)
(/Species: Vespa orientalis (oriental hornet)
(/Species: Vespa orientalis (oriental hornet)
(/Date: 29-Jan-1993 #sequence_revision 29-Jan-1993 #text_change 09-Jul-2004
(/Accession: S10920
R;Tuichibaev, M.U.; Akhmedova, N.U.; Kazakov, I.; Korneev, A.S.; Gagel'gans
Blochemistry (N.Y.) 53, 183-190, 1988
A;Title: Low-molecular-weight peptides of venom of the giant hornet Vespa o
A;Reference number: S06445
A;Accession: S10920
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S10920
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R;Sun, S.C.; Joris, L.; Tsolas, O.
Arch. Biochem. Biophys. 178, 69-78, 1977
A;Title: Purification and crystallization of transaldolase isozyme I and A;Reference number: A12872; MUID:77110646; PMID:556924
A;Accession: A12872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transaldolase (EC 2.2.1.2) I - yeast (Pichia jadinii) (fragment) C;Species: Pichia jadinii, Candida utilis C;Date: 05-un-1987 #sequence_revision 05-Jun-1987 #text_change 09-Jul-2004 C;Accession: Al2872
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C;Date: 20-Uun-2000 #sequence_revision 20-Jun-2000 #text_change 16-Aug-2004
C;Accession: A03188
R;Lote, C.J.; Weiss, J.B.
Biochem. J. 123, 25P, 1971
A;Title: Identification in urine of a low-molecular-weight polar glycopeptide containing A;Reference number: A03188; MUID:72062338; PMID:5126885
A;Molecule type: protein
A;Residues: 1-8 <LOT>
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C;Keywords: transferase
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A;Molecule type: protein
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Pred. No. 2.8e+05;
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0. 2.8e+05;
0;
                                                                                       Korneev, A.S.; Gagel'gans, A.I.
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                                                    giant hornet Vespa orientalis.
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T-cell receptor beta chain V-D-J region (clone 13) (Species: Rattus norvegicus (Norway rat) C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 C;Accession: PH0942 R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vande
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          trypsin (EC 3.4.21.4) precursor - minke whale (fragment) c;Species: Balaenoptera acutorostrata (minke whale, lesser rorqual) C;Date: 17-Jul-1994 #sequence_revision 17-Jul-1994 #text_change 09-Jul-2004 C;Accession: A61328
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C;Keywords: hydrolase; protein digestion; serine proteinase;
F;1-8/Domain: activation peptide #status experimental <APT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Jones, C.M.; Prince, C.A.; Williams, J.S.
Exp. Hematol. 19, 704-709, 1991
A;Title: Purification and amino acid analysis of a human macrophage cytotoxicity-induci A;Reference number: A60427; MUID:91372335; PMID:1909970
A;Accession: A60427
                                                                                                  PH0942
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A; Residues: 1-8 < BRI>
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C;Species: Homo sapiens (man)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Reference number: A61328; MUID:75146765; PMID:1125273
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A;Cross-references: UNIPROT:Q7M471; UNIPARC:UPI000017BF07
C;Keywords: venom
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Local Similarity 28.6%;
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Vandenbark,

A.A.; Wilson, D.B.

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J. Exp. Med. 174, 1467-1476, 1991
A,Fittle: Analysis of T cell receptor beta chains in Lewis rats with experimental allergi A;Reference number: PH0891; MUID:92078857; PMID:1836012
A;Rocession: PH0942
A;Molecule type: mRNA
A;Residues: 1-9 <GOL>
A,Cross-references: UNIPARC:UDI000017C9DD
A;Experimental source: complete Freund's adjuvant-immunized lymph node
A;Rote: the authors translated the codon TGC for residue 2 as Ala
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                       RESULT 15
140697
blotin A - Citrobacter freundii (fragment)
C;Species: Citrobacter freundii
C;Species: Citrobacter freundii
C;Species: Citrobacter freundii
C;Species: Citrobacter freundii
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 09-Jul-2004
C;Accession: 140697
R;Schiuan, D; Campbell, A.
Gene 67, 203-211, 1988
A;Title: Transcriptional regulation and gene arrangement of Escherichia coli, Citrobacte
A;Reference number: 140697; MUID:89006280; PMID:2971595
A;Accession: 140697
A;Accession: 140697
A;Bratus: preliminary; translated from GB/EMBL/DDBJ
A;Residues: 1-4 <RES>
A;Residues: 1-4 <RES>
A;Cross-references: UNIPROT:P13071; UNIPARC:UPI000017AA21; GB:M21922; NID:g144434
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                                                                                                                                                                                                                              Query Match 26.7%; Score 12; DB 2; Length 4; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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Q28742 PABIT
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Q99182 9SMEG
Q726G0 HUMAN
FAR8 MACRS
Q70SM2 HUMAN
Q4X981 PLACH
Q8LPTS MAIZE
P82568 STRPY
Q15900 HUMAN
TAL1 PĪCJA
Q7M4RS HUMAN
Q7M471 VESOR
Q9FSZ2_CICAR
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Q8C339 RAT
Q9QZA8 MOUSE
Q6Q7G0 RANRI
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Q47556 escherichia
Q28742 oryctolagus
Q7m3s5 trypanosoma
O99182 gnatholebia
 Q938r0
P17440
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Q9f8z2
Q8cg39
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1 homo sapian
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RESULT 2
Q47556_ECOLI PRELIMINARY;
ID Q47556;
AC Q47556;
DT 01-NOV-1996 (TrEMBLrel. 01, C:
DT 01-NOV-1996 (TrEMBLrel. 08, L:
DT 01-NOV-1996 (TrEMBLrel. 08, L:
GN Name=pyrI;

Created)
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Last annotation update)
regulatory chain (Fragment).

PRT;

9 AA

g Qy	Qu. Be:	SQ KW	3888	RR	8 R R S	R R R	RA RA	R R R	R RN	288 188		RESULT Q7R8X5 ID Q7								
	Query Match Best Local Matches	EMBL; AABLO1 Hypothetical SEQUENCE 9	Nature -!- CAU EMB	"Genome parasite	van Lin L.H., Jan Salzberg S.L., V	Cho J.K., Q	Silva J.C., Er Peterson J.D.,	STRAIN=17XNL; MEDLINE=22255706; Carlton J.M., Ang	NUCLEOTIDE	Plasmodium Eukaryota;	01-MAR-2004 (01-MAR-2004 (01-MAR-2004 (01-MAR-2004 (Hypothetical	1 PLAY 7R8X5 7R8X5		44	42					
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	; Score 19; DB 2; Length 9; ; Pred. No. 2.2e+06; 3; Mismatches 0; Indels	52.1; -; Genomic_DNA. 4684D447244441E7 CRC64;	here is derived from me shotgun (WGS) entr	rative analysis of the model yoelii.";	rs A.P., Smith H.O., aser C.M., Hoffman S.	gah M., Shoaibi ine J.D., Sinden	Silva J.C., Ermolaeva M.D., Allen J.E., Selengut J.D., Peterson J.D., Pop M., Kosack D.S., Shumway M.F., Biland Reference of the control of t	PubMed=12368865; DOI=10.1038/nature01099 uoli S.V., Suh B.B., Kooij T.W., Pertea		oelii. a; Apicomplexa; Haemosporida; Plas	Created) Last sequence update) Last annotation update)	PRT; 9 AA.	ALIGNMENTS		Q56ST3_VIBOP					Q7M390_BALA
	9; els 0; Gaps 0;		an Y which is	l rodent malaria	White O.R., L., Gardner M.J.,	A., Cummings L.M., R.E., Harris M.A.,	well S.L.,	re01099; Pertea M.,		Plasmodium.				viburnum viburnum	Q56st1 viburnum tr Q56st3 viburnum op	viburnum	malus don zea mays		bacterion glycine m	Q7m390 balaenopter

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RESULT 3
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"Genes encoding Escherichia
pyrB-pyrI operon.";
proc. Natl. Acad. Sci. U.S.A
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Bacteria; Proteobacteria; Gammaproteobacteria;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Alpha-myosin heavy chain (Fragment).
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Lagomorpha; Leporida
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Hoover T.A., Roof W.D., Foltermann
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MEDLINE=82275057; PubMed=7051000;
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MEDLINE=84221901; PubMed=6328491;
Umeda P.K., Sinha
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Proc. Natl. Acad. Sci. U.S.A. 81:3044-3048(1984).
EMBL; K01698; AAA31415.1; -; Genomic_DNA.
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Rabinowitz M.;
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Q7Z6GO_HUMAN PRELIMINARY;
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Q1-OCT-Z003 (TrEMBLrel. 25, Las
Q1-OCT-2003 (TrEMBLrel. 25, Las
Q1-OCT-2003 (TrEMBLrel. 25, Las
Tumarate hydratase (Fragment).
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01-MAR-2004 (TrEMBLrel. 26, Last annotation
phosphoenolpyruvate carboxykinase.
Trypanosoma brucei.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Purification and characterization of phosphoenolpyruvate carboxykinase from Trypanosoma brucei."; Biochim. Biophys. Acta 1249:15-22(1995). PIR; S55696; S55696.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=5691;
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Hunt M., Koehler P
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NON_TER 1
SEQUENCE 7 AA
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01-MAY 1999 (TrEMBLrel. 10, Last
01-JUN 2003 (TrEMBLrel. 24, Last
Cytochrome oxidase I (Fragment).
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                                                                                                                                                                                                                                                                                                                        MEDLINE=20072928; PubMed=10603257; DOI=10.1006/mpev.1999.0656; Murphy W.J., Thomerson J.E., Collier G.E.; "Phylogeny of the Neotropical killifish family Rivulidae"
                                                                                                                                                                                                                                                                                                                                                                                                                                               Mitochondrion.
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GO; GO:0005739; C:mitochondrion;
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 sapiens (Human).
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33.3%;
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RESULT 7

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Wei M.-H., Nickerson M.L., Toro J.R.;
Submitted (MAY-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AY299638; AAP57532.1; -; Genomic_DNA.
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28-FEB-2003 (Rel. 41, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
05-JUL-2004 (Rel. 44, Last annotation update)
FMRFamide-like neuropeptide FLP8 (VSHNNFLRF-amide).
Macrobrachium rosenbergii (Giant fresh water prawn).
Bukaryota, Metazoa, Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Plecoyemata; Caridea
Palaemonoidea; Palaemonidae; Macrobrachium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sithigorngul P., Saraithongkum W., Longyant S., Panchan N., Sithigorngul W., Petsom A.; Sithigorngul W., Petsom A.; "Three more novel FMRFamide-11ke neuropeptide sequences from the eyestalk of the giant freshwater prawn Macrobrachium rosenbergii Peptides 22:191-197(2001).
                                                                                                                                                                                                                                                                                                                                                                          Amidation; Direct MOD_RES 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=21107394; PubMed=11179812; DOI=10.1016/S0196-9781(00)00382-X; Sithigorngul P., Saraithongkum W., Longyant S., Panchan N.,
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Mammalia; Eutheria;
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les 3; Conserv
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MASS SPECTROMETRY: MW=1133.8; METHOD=MALDI; RANGE=1-9; NOTE=Ref.1.

SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
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Nucleof Serviman M., Raine J.D., Carlton J.M., Kooij T.W.A.,

Berriman M., Florens L., Janssen C.S., Pain A., Christophides G.K.,

James K., Rutherford K., Harris B., Harris D., Churcher C.,

Quail M.A., Ormond D., Doggett J., Trueman H.E., Mendoza J.,

Bidwell S.L., Rajandream M.A., Carucci D.J, Yates J.R., Kafatos F.C.,

Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S.;

"A comprehensive survey of the Plasmodium life cycle by genomic,

transcriptomic, and proteomic analyses.";

Science 307:82-86 (2005).

-!- CAUTION: The sequence shown here is derived from an

FMRI./GenBank/DDBJ whole genome shotgun (WGS) entry which is
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Eukaryota; Alveolata;
NCBI_TaxID=5825;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-SEP-2005 (TrEMBLrel. 31, Created)
13-SEP-2005 (TrEMBLrel. 31, Last sequence up
13-SEP-2005 (TrEMBLrel. 31, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NON TER
                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                                   Hypothetical protein.
NON_TER 1
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Q4X981;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Mammary gland;
Pinte S., Guerardel C., Deltour S., Godwin A.K., Lepring
"Identification of a second G-C-rich promoter conserved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (Human).
Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27, 05-JUL-2004 (TrEMBLrel. 27, Hypermethylated in cancer 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     murine and rat tumor suppressor genes HIC1."; Oncogene 23:4023-4031(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                               preliminary data.
; CAAJ01008648; CAH86543.1;
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                                                               \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 DLLEA 9
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                                                                                                                                                                     Similarity
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3; Conserv
                                                        ATHEET 6
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                                                                                                                                       Conservative
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1010 MW;
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16.7%;
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                                                                                                                               Score 16; DB 2;
Pred. No. 2.2e+06;
; Mismatches 0
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                               -; Genomic_DNA.
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                                                                                                                                   0,
                                                                                                                                                                                                   Length 9;
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RESULT 10
QBLPT5_MAIZE PRELIMINARY;
ID QBLPT5_MAIZE PRELIMINARY;

PRT;

9

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PRESULT 11
PREZS68 STRPY
PREZS68 STRPY
PREZS68 STRPY
PREZS68;

AC
PREZS68;

DT
01-OCT-2000 (TrEMBLrel. 15, 10)
DT
01-OCT-2003 (TrEMBLrel. 25, 10)
DT
01-OCT-
                                                   RESULT 12
Q15900 HUMAN
Q15900 HUMAN PRELIMINARY;
ID Q15900;
AC Q15900;
DT 01-NOV-1996 (TremBLrel. 01, Cr
DT 01-NOV-1996 (TremBLrel. 19, La
DT 01-DEC-2001 (TremBLrel. 19, La
DT 01-DEC-101 (TremBLrel. 19, La
DEC. Homo sapiens (clone XP7B11A)
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Best Local S
Matches 2
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Best Local
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01-OCT-2002
01-OCT-2002
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01-OCT-2000 (TERMBLICE. 15, La
01-OCT-2003 (TERMBLICE. 25, L
01-OCT-2003 (TERMBLICE). 25, L
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Morgante M., Rafalski J.A.;
Submitted (APR-2002) to the EMBL/GenBar
EMBL; AY094310; AAM21836.1; -; Genomic
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Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; E
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2; Conserv
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(TrEMBLrel. 22,
(TrEMBLrel. 22,
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                                                                                                                                                                                                                                                                                                                                                          Conservative
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  Euarchontoglires; Primates;
                       Chordata;
                                                                                                                                                                                                                                                                                                                                                                             35.6%;
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. 15, Last sequence upo
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D-page (Fragment).
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Pred. No.
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Pred. No.
      Craniata; Vertebrata; Euteleostomi;
oglires; Primates; Catarrhini; Homin
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RESULT 13
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Hum. Mol. Genet. 0:0-0(1995)
EMBL; L32079; AAA73890.1; -;
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Coolbaugh M.I., Chinault C.A., Baldini
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Placenta;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Berriman M., Florens L., Janssen C.S., Pain A., Christophides G. James K., Rutherford K., Harris B., Harris D., Churcher C., James K., Ormond D., Doggett J., Trueman H.E., Mendoza J., Quail M.A., Ormond D., Doggett J., Tuchen H.E., Mendoza J., Bidwell S.L., Rajandream M.A., Carucci D.J., Yates J.R., Kafatos Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S., Janse C.J., Barrell B., Turner C.M.R., Waters A.P., Sinden R.S., "A comprehensive survey of the Plasmodium life cycle by genomic,"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Plasmodium chabaudi.
Bukaryota; Alveolata; Apicomplexa;
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EMBL; CAAJ01003387; CAH79935.1; -; Genomic_DNA.
                                              01-NOV-1999 (TrEMBLrel.
01-NOV-1999 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transcriptomic, and proteomic analyses."; Science 307:82-86(2005).
                                                                                                               Q9XJN0 9VIRU PRELIMINARY;
      Bacteriophage phi-10
                         P10 (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAUTION: The sequence shown here is derived EMBL/GenBank/DDBJ whole genome shotgun (WGS)
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Karras M., Raine J.D., Carlton J.M., Kooij T.W.A.,
M. Florens L., Janssen C.S., Pain A., Christophides
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MINDICINE-9350412; PubMed=10419946;
Mindich L., Qiao X., Qiao J., Onodera S., Romantschuk M.,
Hoogstraten D.;
"Isolation of additional bacteriophages with genomes of segmented double-stranded RNA.";
J. Bacteriol. 181:4505-4508(1999).
EMUEL; AR125675; AAD225555.1; -; Genomic_RNA.
NON_TER 9
                                                                                                                                                                                                          Tryptophan biosynthesis. NON_TER 6 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This Swiss-Prot entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TRPI_PSEPU STANDARD; PRT; 6 AA. 1936414; 01-JUN-1994 (Rel. 29, Created) 01-JUN-1994 (Rel. 29, Last sequence update) 13-SEP-2005 (Rel. 48, Last annotation update) HTH-type transcriptional regulator trpi (TrpBA)
                                                                                                                                                                                                                                                   InterPro; IPR000847; HTH_LysR.

PROSITE; PS50931; HTH_LYSR; PARTIAL.

Activator; Amino-acid biosynthesis; Aromatic amino acid biosynthesis;

DNA-binding; Transcription; Transcription regulation;
                                                                                                                                                                                                                                                                                                                                                                  EMBL; X13299; CAA31660.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-89335826; PubMed-2503057; DOI=10.1016/0300-9084(89)90183-1; Eberly L., Crawford I.P.; "DNA sequence of the tryptophan synthase genes of Pseudomonas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pseudomonas putida.
Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
Pseudomonadaceae; Pseudomonas.
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                                                              Local Similarity
nes 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION: Activates the expression of the trpBA genes encoding to two tryptophan synthase subunits. In the absence of the inducer (indoleglycerol phosphate), trpl binds upstream of the trpAB operon, overlapping its own promoter region.

SIMILARITY: Contains 1 HTH lysR-type DNA-binding domain.
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                                                         31.1%; Score 14; DB 1; Le ilarity 100.0%; Pred. No. 2.2e+06; Conservative 0; Mismatches 0;
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Listing first 45 summaries
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/AaCkfiles1.pep:*
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US-09-269-250R-3
US-09-269-250R-3
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US-08-159-339A-370
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US-08-428-823-9
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TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:

Fragments

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RESULT 1 US-09-269-250E-20 Sequence 20, Applica Patent No. 6830883 GENERAL INFORMATION: APPLICANT: GOULINY, APPLICANT: HOUSEN COLORER FITTE OF INVENTION: FITTE REFERENCE: 599 CURRENT APPLICATION CURRENT FILING DATE NUMBER OF SEQ ID NO SOFTWARE: Patentin SEQ ID NO 20 LENGTH: 9 FYDER, DET		44444333333333333333333333333333333333	22
ULT 1 09-269-250E-20 equence 20, Applic atent No. 6830883 ENERAL INFORMATION APPLICANT: GOLLINY, TITLE OF INVENTION FILE REFERENCE: 58 CURRENT APPLICATION CURRENT APPLICATION CURRENT FILING DAT NUMBER OF SEQ ID N SOFTMARE: Patentin EQ ID NO 20 TENCTH: 99 TYPE. BET		222222222222222222222222222222222222222	2 2
SULT 1 -09-269-250E-20 Sequence 20, Application US/09269250E Sequence 20, Application US/09269250E Sequence 10, 6830883 GENERAL INFORMATION: APPLICANT: GOULMY, Elsa TITLE OF INVENTION: METHOD FOR TYPIN TILE REFERENCE: 58994 CURRENT FILING DATE: 1999-05-21 RUMBER OF SEQ ID NOS: 38 SOFTWARE: Patentin version 3.1 SEQ ID NO 20 TYPE. DET		**************************************	
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G OF MINOR 9,250E	ALIGNMENTS	US-08-459-568-70 US-08-399-411-20 US-08-399-411-20 US-08-39-566-472-20 US-09-586-472-20 US-09-586-472-20 US-09-586-472-75 US-08-399-411-75 US-09-419-826-7 US-09-419-826-7 US-09-528-706-75 US-08-37-30 US-09-528-706-75 US-08-343-602-2 US-08-459-568-72 US-08-459-568-72	US-09-217-609A-11
HISTOCOMPATIBILITY ANTIGEN HA-1		Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence	Sequence
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Sequence 2, Application US/09489760
Patent No. 6678375
GENERAL INFORMATION:
APPLICANT: RIJKSUMIVERSITEIT TE Leidin
APPLICANT: ROULMY, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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LENGTH: 9
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Pred. No. 4.6e+05;
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APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEO ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 5
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RESULT 5 '
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CURRENT FILING DATE: 199-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 18
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
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Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
FILE REFERENCE: 58994
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Best Local Similarity
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Best Local
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                                                                                                                                                                                             TYPE: PRT ORGANISM: KIAA0223 partial complementary DNA
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
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US-09-489-760-4
                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO
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Best Local S
Matches 8
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GENERAL INFORMATION:
APPLICANT: GOULT, Elsa
APPLICANT: GOULT, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 29, Application US/09269250E Patent No. 6830883
                                                                                                                                   Best Loc
Matches
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                                                                                                                                                     Query Match
Best Local
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APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                      LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: MISC_FEATURE LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a histidine (H) or an arginine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Human
                                                                                                                                                                                                                                                                  ORGANISM: Histocompatibility antigen FEATURE: NAME/KEY: PEPTIDE
                                                                                                                                                                                                              LOCATION: (3)
OTHER INFORMATION: AMINO ACID X REPRESENTS A HISTIDINE OR AN ARGININE OTHER INFORMATION: RESIDUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 9
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8; Conserv
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                                                                   VLXDDLLEA
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88.9%;
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Pred. No. 4.6e+05;
0; Mismatches 1
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Pred. No.
                                                                                                                                         ed. No. 4.6e+05;
Mismatches 1
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Sequence 4, Application US/09489760 Patent No. 6878375

GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin

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COCATION: (2)..(7)
OTHER INFORMATION: AMINO ACIDS X REPRESENT LEUCINE OR ISOLEUCINE
OTHER INFORMATION: RESIDUES
US-09-489-760-4
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Best Local Similarity
""" hes 6; Conserv
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GENERAL INFORMATION:
APPLICANT: GOULMY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
                                                                                                                                                                      Sequence 5, Application US/09025819 Patent No. 6225097
                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Obata, Shusei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
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APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 9
TYPE: PRT
ORGANISM: Histocompatibility antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: HUMAN
                                      APPLICANT: Obata, Shusei
APPLICANT: Nishino, Tokuzo
APPLICANT: Koyama, Tanetoshi
APPLICANT: Sato, Yoshihiro
TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH: 8
                                 CORRESPONDENCE ADDRESS
   ADDRESSEE:
STREET: 15
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8: KENYON & KENYON
1500 K Street, N.W
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88.9%;
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Pred. No.
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Pred. No. 4.6e+05;
0; Mismatches 0;
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4.6e+05;
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RESULT 10
US-09-808-126-5
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Best Local Similarity
Matches 4; Conserv
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ETLING DATE: 17-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: Khalilian, Houri
REGISTRATION NUMBER: 39,546
REFERENCE/DOCKET NUMBER: 10235/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: UFILING DATE: 19-FEB-1
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                         ZIP: 20005

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/808,126
FILING DATE: 08-May-2001
CLASSIFICATION: <Unknown>
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STRANDEDNESS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/025,819
FILING DATE: <Unknown>
ATTORNEY/AGENT INPORMATION:
NAME: Khalilian, Houri
REGISTRATION NUMBER: 39,546
                                                                                                                                                                                                                                                                                                                                                                                                         Sato, Yoshihiro
TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Obata, Shusei
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STREET: 1500 K Street, N.W.
CITY: Washington
STATE: DC
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Koyama, Tanetoshi
Sato, Yoshihiro
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Pred. No. 4.6e+05;
1; Mismatches 0
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RESULT 11
US-09-803-951-5
(Sequence 5, Application
Patent No. 6413761
GENERAL INFORMATION:
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                                                                  ; TOPOLECULE TYPE: POPTIGE
; MOLECULE TYPE: Poptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 5:
US-09-803-951-5
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Matches 4; Conservative
Quary Match
Best Local Similarity 80.4
Matches 4; Conservative
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INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                    TELEFAX: 202-220-4201
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                          APPLICATION NUMBER: 09/025,819
FILING DATE: «Unknown)
ATTORNEY/AGENT INFORMATION:
NAME: Khalilian, Houri
REGISTRATION NUMBER: 39,546
REFERENCE/DOCKET NUMBER: 10235/2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-220-4200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 10235/2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-220-4200
                                                                                                                                                                                                                                                                                                                                                                                                              SOPTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/803,951
FILING DATE: 13-Mar-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: KENYON & KENYON
STREET: 1500 K Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nishino, Tokuzo
Koyama, Tannetoshi
Sato, Yoshihiro
TITLE OF INVENTION: DECAPRENYL DIPHOSPHATE SYNTHETASE GENE
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Obata, Shusei
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STRANDEDNESS: <Unknown>
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                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Washington
                                                                                                                                                                                          LENGTH: 7 amino acids
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80.0%;
                   55.6%;
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Pred. No. 4.6e+05;
   Score 25; DB 2; Length 7; Pred. No. 4.6e+05; 1; Mismatches 0; Indels
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US-09-305-923A-7
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GENERAL INFORMATION:
APPLICANT: OSCANIN, Kirill
APPLICANT: SILVERMAN, LAUREN
TITLE OF INVENTION: YEAST CELLS HAVING MUTATIONS IN Stp22 AND USES THEREFOR TITLE REFERENCE: CPI-091
CURRENT APPLICATION NUMBER: US/09/305,923A
CURRENT APPLICATION NUMBER: 60/084,420
PRIOR APPLICATION NUMBER: 60/084,420
PRIOR APPLICATION NUMBER: 60/084,420
PRIOR FILING DATE: 1998-05-06
NUMBER OF SEQ ID NOS: 9
SOFTWARE: PATENTIN Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                Sequence 7, Application US/09305923A Patent No. 6355473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Klein, Christine A.
APPLICANT: Murphy, Andrew J. M.
TITLE OF INVENTION: Methods and
TITLE OF INVENTION: Identifying
NUMBER OF SEQUENCES: 98
                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Catherine J. Kara
REGISTRATION NUMBER: 41,106
REFERENCE/DOCKET NUMBER: CPI-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (617)227-740
TELEFAX: (617)227-4214
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ABCII(text)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/582,333A
PILING DATE: 17-JAN-1996
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 28 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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CITY: Boston
STATE: Massac
                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 9 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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33.3%;
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Pred. No. 4.6e+05;
5; Mismatches 1
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APPLICANT: FOWNIES, DANA M.

APPLICANT: FOWNIES, DANA M.

TITLE OF INVENTION: YEAST CELLS EXPRESSING MODIFIED G PROTEINS AND METHODS

FILE REFERENCE: CPI-012CP8

CURRENT APPLICATION NUMBER: US/08/946,298A

CURRENT FILING DATE: 1997-10-07

EARLIER APPLICATION NUMBER: 08/689,172

EARLIER APPLICATION NUMBER: 08/689,172

EARLIER FILING DATE: 1996-08-06

EARLIER FILING DATE: 1996-08-06

EARLIER FILING DATE: 1996-08-11

EARLIER FILING DATE: 1996-08-05

EARLIER APPLICATION NUMBER: 08/463,181

EARLIER APPLICATION NUMBER: 08/322,137

EARLIER APPLICATION NUMBER: 08/302,137

EARLIER APPLICATION NUMBER: 08/302,137

EARLIER APPLICATION NUMBER: 08/303,33

EARLIER FILING DATE: 1994-10-13

EARLIER FILING DATE: 1994-01-31

EARLIER FILING DATE: 1994-01-31

EARLIER APPLICATION NUMBER: 08/190,328

EARLIER APPLICATI
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US-09-217-609A-8
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; TYPB: PRT
; ORGANISM: Saccharomyces cerevisiae
US-08-946-298-4
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US-08-946-298-4
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; TYPE: PRT
; ORGANIEM: Saccharomyces cerevisiae
US-09-305-923A-7
                                                                                                                                                                                                                                Patent No. 6071733

GENERAL INFORMATION:

APPLICANT: MURAMATS

APPLICANT: KOIKE, J

APPLICANT: CGURA, H

APPLICANT: KOYAMA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 33.3
Marches 3; Conservative
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Best Local Similarity 33.3%;
Matches 3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4, Application US/08946298A Patent No. 6864060
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                                                                  APPLICANT: SHIMIZÜ, Naoto
APPLICANT: CHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Kenyon & Kenyon
STREET: 1025 Connecticut Avenue, NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 VLHDDLLEA 9
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1 LIHEDIAKA 9
                                                                                                                                                             MURAMATSU, Masayoshi
KOIKE, Ayumi
OGURA, KYOZO
KOYAMA, Tanetoshi
SHIMIZU, Naoto
CHO, Yenwin
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33.3%;
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
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          Suite 600
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Search completed: April Job time : 28.6667 secs
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/873,23
FILING DATE: 11-Jun-1997
ATTORNEY/AGENT INPORMATION:
NAME: TOPPENETIT, Judith L.
REGISTRATION NUMBER: 39,048
REFERENCE/DOCKET NUMBER: 1023.
TELECOMMUNICATION INPORMATION:
TELEPAN: 202-429-1776
TELEPAN: 202-429-0796
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                      Matches
                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: DC
COUNTRY: US
ZIP: 20036
ZIP: 20036
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
SOFTWARE: WordPerfect 6.1 for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/217,609A
FILING DATE:
CLASSITED: ATTOM.
                                                                                                                                                                                                                                            TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                        LENGTH: 6 amino acids
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Pred. No. 4.6e+05;
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Gaps

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Minimum DB
Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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   seq length: 0 seq length: 9
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Match Length
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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Gapop 10.0 , Gapext 0.5
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US-10-623-176-2
US-10-971-217-2
US-10-971-217-2
US-11-007-740-20
US-10-623-176-41
US-10-623-176-40
US-10-623-176-40
US-10-623-176-10
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20, Appl
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16, Appl
17, Appli
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2 9nce icat icat LICA LICA LICA	р р	Query Match Best Local Matches	TYPE: 9 TYPE: PRT ORGANISM: FEATURE: OTHER INF FEATURE: NAME/KEY: LOCATION: 0-623-176	PRIOR APPL PRIOR FILI NUMBER OF SOFTWARE: SEQ ID NO 2	RIOR FILI RIOR APPI	URRENT AI	APPLICANT: APPLICANT: APPLICANT: APPLICANT: TITLE OF IN	1 623- ence icat		C B C C C C C C C C C C C C C C C C C C	ω4υ	N H O O Ø	
217-2 2, Application No. US20 ion No. US20 ION TO COLUMN IN FORMATION: ANT: Goulmy, ANT: Hunt, D ANT: Engelha	VLHDDLLEA	Simi 9;	Artif ORMATI SITE (1)	CATIC G DAI EQ II atent	ICATIO	PLICAT	Goulm Hunt Enge VENTI	<i>_</i> '		22 4 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5			
ion 0401 Elsa Enal	LLEA 9	vat	ρ	TUMB 200 Ver	N NUMBER E: 1997- N NUMBER E: 1998-	ENT APPLICATION NUMBER: US/ ENT FILING DATE: 2003-07-11 A APPLICATION NUMBER: 09/48 R FILING DATE: 2000-01-21	Els onal rd, HA-	ion 0400		ம் ம் ம் க் க் க் க் க் க்			
US/1079 [91268A1 [9126.M A.J.M d F Victor		100.0 100.0	l Sequence Description	ER: JP 0-01-2 77 . 2.1	R: EP '-07-23 R: PCT -07-23	BER:	A.J.M. ld F. Victor 1 epit	US/1062)92446A1		υυυυυυ <i>ι</i> αα ωυ4αυυ4ου			
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		Score 45; Pred. No. Mismatc	Artif	504165	03.0	, 17	d uses		ALIGNMENTS	09-747-774N 09-953-354- 10-953-354- 110-267-074- 110-267-078- 110-967-087- 110-967-087- 110-808-187- 110-807-807-	825- 166- 166- 803-	705- 166- 166- 166-	
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		B 4; 7e+06					thereo			1 & 4 & & & & & & & & & & & & & & & & &	48 -161 -162 5	10-705-459-280 10-166-225A-163 10-166-225A-164 10-166-225A-165 10-166-225A-166	
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                                                                                                                                                                                                          RESULT 4
US-11-007-740-20
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Matches
               Sequence 20, Application US/11007740
publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
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Best Local Similarity
Matches 9; Conserv
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SEQ ID NO 1
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: Patentin Ver. 2.1
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CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
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CURRENT FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: PCT/NL02/00791
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR PILING DATE: 2001-12-05
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
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TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunot
TITLE OF INVENTION: of tumors
FILE REFERENCE: 2183-6479US
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ORGANISM: Artificial sequence
FEATURE:
PRIOR FILING DATE: 1999-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
OTHER INFORMATION: HA-1 peptide
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; OTHER INFORMATION: Description of Artificial Sequence: Exon Fragments US-11-007-740-20
RESULT 6
US-10-623-176-45
Sequence 45, Application US/10623176
Publication No. US20040092446A1
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR PPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses
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CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
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TYPE: PRT
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nes 8; Conserv
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100.0%; Pred. No.
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Pred. No. 1.7e+06;
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CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18

APPLICATION NUMBER: 09/489,760 FILING DATE: 2000-01-21

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; FEATURE:
; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-45
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PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: BP 97202303.0
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: D 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 10
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US-10-623-176-10
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APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILS REFERENCE: 2183-6047US
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SOFTWARE: PatentIn Ver. 2.1
SQ ID NO 45
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CURRENT FILING DATE: 2003-07-18
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PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
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TYPE: PRT
ORGANISM: Artificial Sequence
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No. US20040092446A1
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RESULT 8

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Query Match
Best Local Similarity
""" hes 7; Conserv?
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APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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                                                                                                                                           ; OTHER INFORMATION: obtained from KIAA0223 partial complementary US-10-791-217-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 9
US-10-791-217-5
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                                                                                                                                                                                                                                                                                                                                 APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
ITILE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
                                                           Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                       SEQ ID NO 5
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/10791217 Publication No. US20040191268A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: SITE LOCATION: (1)...
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OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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1 VLHDDLLEA 9
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8; Conser
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No. US20040092446A1
                                                             Conservative
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100.0%; Pr
                                                                               82.2%;
88.9%;
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Pred. No. 1.7e+06;
                                                             Score 37; DB 4; Le
Pred. No. 1.7e+06;
0; Mismatches 1;
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                     Query Match
Best Local Similarity
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APPLICANT: GOULMY, EISA
APPLICANT: GOULMY, METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18, Appublication No.
                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT EPPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                      FEATURE:
NAME/KEY: SITE
LOCATION: (1)..(9)
                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide OTHER INFORMATION: wherein X can be R or H
                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                                                                                           ENGTH: 9
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les 8; Conserv
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    1 VLHDDLLEA 9
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                                          Conservative
                                                             80.0%;
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88.9%; Pred. No. 1.70
tive 0; Mismatches
                                        0,
                                                           Score 36; DB 4;
Pred. No. 1.7e+06;
                                            Mismatches
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Sequence 1, Application US/10791217

Publication No. US20040191268A1

GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
ITTLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US

CURRENT APPLICATION NUMBER: US/10/791,217

CURRENT FILING DATE: 2004-03-02

PRIOR APPLICATION NUMBER: US/09/489,760

PRIOR FILING DATE: 1908-07-23

NUMBER OF SEQ ID NOS: 17

COPPENSED: DATE: 1998-07-23

NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: GOULTNY, E18 A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 1090-01-21
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EP7/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-01-24
NUMBER: DETONERE: US/1000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER: DETONERE: DETONERE: US/1000-504165
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US-10-791-217-1
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Publication No. US20040092446A1
GENERAL INFORMATION:
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SEQ ID NO 40
                                                                                                                                                                                            SEQ ID NO
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                                                                                                                                           LENGTH: 9
TYPE: PRT
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FEATURE: OF THE INFORMATION: obtained from histocompatibility antigen FEATURE: NAME/KEY: MISC FEATURE LOCATION: (3)...(3)
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                                                                                                                     ORGANISM: Artificial sequence
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100.0%; Pred. No. 1.
cive 0; Mismatches
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APPLICANT: HOULDY, E1s A.J.M.

APPLICANT: Hunt, Donald F.

APPLICANT: Engelhard, Victor H.

PRICE OF INVENTION: HA-1 epitopes and uses thereof

FILE REFERENCE: 2183-6047US

CURRENT APPLICATION NUMBER: US/10/623,176

CURRENT APPLICATION NUMBER: 09/489,760

PRIOR APPLICATION NUMBER: E97202303.0

PRIOR APPLICATION NUMBER: E97202303.0

PRIOR APPLICATION NUMBER: E9700424

PRIOR APPLICATION NUMBER: DON-01-21

PRIOR APPLICATION NUMBER: DON-01-24

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR PILING DATE: 1308-07-23

PRIOR APPLICATION NUMBER: JP 2000-504165

PRIOR PILING DATE: 2000-01-24

NUMBER OF SEQ ID NOS: 77

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 42

LENGTH: 9

TYPE: PRT
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US-10-623-176-42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 14
US-11-007-740-29
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Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR PILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PatentIn version 3.1
SEQ ID NO 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 42, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:
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Best Local Similarity
Matches 8; Conservat
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TYPE: PRT
ORGANISM: Human
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
                              ORGANISM: Artificial Sequence FEATURE:
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Pred. No. 1.7e+06;
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 9
   Published Applications AA New:*

1: /SIDSS/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

2: /SIDSS/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/US03_NEW_PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

7: /SIDSS/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

8: /SIDSS/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
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  BLOSUM62
Gapop 10.0 ,
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-11-010-748A-11
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US-11-129-143-166
US-11-129-143-162
US-11-034-399-1155
US-10-485-788A-355
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US-11-033-039-785
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          Sequence
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  Sequence
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  11, Appl
163, App
164, App
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166, App
161, App
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353, App
353, App
354, App
352, App
352, App
352, App
352, App
354, App
3715, Ap
5715, Ap
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RESULT 2 US-11-010-748A-12 J Sequence 12, Application US/11010748A PUDIcation No. US20050244421A1 GENERAL INFORMATION: APPLICANT: Merck Patent GmbH APPLICANT: STRITTMATTER, Wolfgang APPLICANT: MOLL, Heidrun APPLICANT: SCHARM, Burkhard	Oy 1 VLHDDLLEA 9 Db 1 VLHDDLLEA 9	Query Match 100.0%; Score 45; DB 7 Best Local Similarity 100.0%; Pred. No. 1.4e+ Matches 9; Conservative 0; Mismatches	; TYPE: PAT ; ORGANISM: Homo sapiens ; PEATURE: ; NAME/KEY: MISC FEATURE ; LOCATION: (1)(9) ; OTHER INFORMATION: Human minor histocompatibili ; OTHER INFORMATION: ell epitopes US-11-010-748A-11	; PRIOR APPLICATION NUMBER: EP02013423.5 ; PRIOR FILING DATE: 2002-06-13 ; NUMBER OF SEQ ID NOS: 926 ; SOFTWARE: PatentIn version 3.1 ; SEQ ID NO 11 ; LENGTH: 9	CURRENT APPLICATION NUMBER: US/11/010 CURRENT FILING DATE: 2004-12-13 PRIOR APPLICATION NUMBER: PCT/EP03/06 PRIOR FILING DATE: 2003-06-13	APPLI APPLI TITLE	ESULT 1 S-11-010-748A-11 S-quence 11, Application US/110 Publication No. US20050244421A1 GENERAL INFORMATION: APPLICANT: Merck Patent GmbH	ALIGNMENTS	20 44.4 8 7 US-11-021 20 44.4 8 7 US-11-021 20 44.4 8 7 US-11-021	20 44.4 6 6 US-10-857 20 44.4 6 6 US-10-857	20 44.4 5 7 US-11-129 20 44.4 5 7 US-11-129 20 44.4 5 6 US-10-857	35 20 44.4 5 7 US-11-129-143-169 36 20 44.4 5 7 US-11-129-143-170 37 20 44.4 5 7 US-11-120-143-170	20 44.4 4 7 US-11-108 20 44.4 4 7 US-11-033 20 44.4 5 7 US-11-129	20 44.4 4 7 US-11-019 20 44.4 4 7 US-11-174	21 46.7 7 6 US-10-510-155-28 20 44.4 4 6 US-10-657-295-3	21 46.7 4 6 US-10-485-788
); Length 9; +05; 0; Indels 0; Gaps 0;	lity antigens HA-1 characterized			OR MODULATING IMMUNE RESPONSE			sequence Sequence	Sequence 397, Sequence 667,	Sequence 173 Sequence 173 Sequence 367	Sequence 169, Sequence 170,	Sequence 69, Sequence 3, Sequence 160	Sequence 5,	Sequence 28 Sequence 26: Sequence 36:	Sequence 351,

by T

SCHARM, Burkhard

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US-11-129-143-164
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APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, W
                                                                                               Sequence 164, Application US/11129143
Publication No. US20050266518A1
GENERAL INFORMATION:
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Best Local (
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SEQ ID NO 163
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TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
          APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
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PRIOR FILING DATE: 2002-06-13
NUMBER OF SEQ ID NOS: 926
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CURRENT FILING DATE: 2004-12-13
PRIOR APPLICATION NUMBER: PCT/EP03/06251
PRIOR FILING DATE: 2003-06-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: MER-136
 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KBY: MISC_FRATURE LOCATION: (1)..(9) COTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by TOTHER INFORMATION: ell epitopes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Bacillus stearothermophilus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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HUMBELIN, Markus
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YELISEEV, Alexei A.
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80.0%;
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Pred. No. 1.4e+05;
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                   SEQ ID NO 166
LENGTH: 6
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APPLICANT:
APPLICANT:
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APPLICANT:
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                                                 CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: PatentIn version 3.1
                                                                                                                     APPLICANT: YELISÉEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
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BRETZEL, Werner HUMBELIN, Markus LOPEZ-ULIBARRI, Rual MAYER, Anne F.

Alan

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; ORGANISM: Bacillus subtilis US-11-129-143-164
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US-11-129-143-165
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SEQ ID NO 165
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CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOFTWARE: PatentIn version 3.1
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APPLICANT:
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CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
                                                                                                                                                                                                                                                                                                                    APPLICANT: BERRY, APPLICANT: BRETZ
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                                                                                                                                                                                                                          APPLICANT: MAYER, Anne F.
APPLICANT: YELISSEY, Alexei A.
APPLICANT: YELISSEY, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: C38435/121966
                                                                                                    LENGTH: 6
TYPE: PRT
ORGANISM: Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
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2 LHDDL 6
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LOPEZ-ULIBARRI, Rual
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5. US20050266518A1
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                                            57.8%;
80.0%;
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80.0%;
                                            Score 26; DB 7;
Pred. No. 1.4e+05;
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Pred. No.
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APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOPTWARE: Patentin version 3.1
SECURIOR ON 162
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Best Local Similarity 80.0
                                                      Matches
                                                                     Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 162, Application US/11129143 Publication No: US20050266518A1
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Best Local Similarity
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SEQ ID NO 161
LENGTH: 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: YELISEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                           LENGTH: 6
TYPE: PRT
                                                                                                                                           ORGANISM: Rhizobium
2 LHDDL 6
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                                                   Similarity 4; Conserv
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HUMBELIN, Markus
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                                                      Conservative
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                                                                                                                                       sp. strain NGR234
                                                                   55.6%;
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80.0%;
                                                 Score 25; DB 7; Le
Pred. No. 1.4e+05;
1; Mismatches 0;

    Mismatches

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Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
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                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR PILING DATE: 2001-08-03
PRIOR PILING DATE: 2001-08-03
PRIOR FILING DATE: 2002-02-25
PRIOR FILING DATE: 2002-09-02
PRIOR FILING DATE: 2002-09-02
NUMBER OF SEQ ID NOS: 841
SOFTWARE: PatentIn version 3.1
SSEQ ID NO 353
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                                                                                                                                                        ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-485-788A-353
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US-11-004-399-1155
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Best Local Similarity 100.
Thes 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn version 3.1 SEQ ID NO 1155
                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schweizer, Johannes
APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions
FILE REFERENCE: 20054-003320US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/11/004,399
CURRENT FILING DATE: 2004-12-03
PRIOR APPLICATION NUMBER: US 60/527,637
PRIOR FILING DATE: 2003-12-03
NUMBER OF SEQ ID NOS: 4043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Ramalingam, Sathiskumar
APPLICANT: Poon, Leo Lit Man
APPLICANT: Peiris, Joseph Sriyal Malik
TITLE OF INVENTION: Genetically Modified Plants Comprising SARS-CoV Viral Nucleotide
TITLE OF INVENTION: Sequences and Methods of Use Thereof For Immunization Against S
FILE REFERENCE: 2587/73166/RDK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Chye, Mee Lee
APPLICANT: Li, Hong Ye
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
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                1 LVHDDV
                                                 1 VLHDDL 6
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3; Conserv
                                                                                    Conservative
                                                                                                    51.1%;
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                                                                                  Score 23; DB 6;
Pred. No. 1.4e+05;
3; Mismatches 0
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US-10-485-788A-354

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; ORGANISM: Homo sapiens
US-10-485-788A-354
                                                                                                                                ; LENGTH: 8
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-485-788A-355
                                                                                                                                                                                                                                                                                                                               APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
FITTLE OF INVENTION: Molecular Interactions in Cells
FILE REFERENCE: 20054-003320US
CURRENT APPLICATION NUMBER: US/10/485,788A
CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR FILING DATE: 2002-02-25
PRIOR FILING DATE: 2002-02-25
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CURRENT FILING DATE: 2004-02-03
PRIOR APPLICATION NUMBER: US 60/309,841
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR FILING DATE: 2002-02-25
PRIOR APPLICATION NUMBER: WD PCT/US02/24655
PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
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APPLICANT: Lu, Peter S.
APPLICANT: Rabinowitz, Joshua D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn version 3.1 SEQ ID NO 354
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Best Local Similarity 50.0
Matches 3; Conservative
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PRIOR FILING DATE: 2002-08-02
NUMBER OF SEQ ID NOS: 841
SOFTWARE: PatentIn version 3.1
SEQ ID NO 355
                                                           Query Match
Best Local Similarity
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APPLICANT:
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APPLICANT: Schweizer, Johannes
APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions in Cells
FILE REFERENCE: 20054-003320US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Lu, Peter S.
APPLICANT: Rabinowitz, Joshua D.
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1 ALHDDL 6
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                                           Conservative
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50.0%;
                                         51.1%; Score 23; DB 6; L6
50.0%; Pred. No. 1.4e+05;
tive 3; Mismatches 0;
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Pred. No. 1.4e+05;
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GENERAL INFORMATION:

APPLICANT: HUMPHREYS, ROBERT

APPLICANT: XU, MINZHEN

TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES

FILE REFERENCE: REH-2017US01

CURRENT APPLICATION NUMBER: US/11/033,039

CURRENT FILING DATE: 2005-01-11

PRIOR APPLICATION NUMBER: 10/245,871

PRIOR APPLICATION NUMBER: 10/197,000

PRIOR APPLICATION NUMBER: 10/197,000

PRIOR APPLICATION NUMBER: 10/197,000

PRIOR APPLICATION NUMBER: 09/396,813

PRIOR APPLICATION NUMBER: 09/396,813

PRIOR APPLICATION NUMBER: 09/396,813

PRIOR PILING DATE: 1999-09-14

NUMBER OF SEG ID NOS: 1452

SOFTWARE: PatentIn version 3.3

SEG ID NO 809
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US-11-033-039-785
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US-11-033-039-785
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US-11-033-039-809
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Publication No. US20060002947A1
GENERAL INFORMATION:
APPLICANT: HUMPHREYS, ROBERT
APPLICANT: XU, MINZHEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn version 3.3 SEQ ID NO 785
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Best Local Similarity 44.4%;
Matches 4; Conservative
                                                                                           Matches
                                                                                                           Query Match
Best Local Similarity 44.4%;
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TITLE OF INVENTION: LI-KEY/ANTIGENIC EPITOPE HYBRID PEPTIDE VACCINES
FILE REFERENCE: REH-2017US01
CURRENT APPLICATION NUMBER: US/11/033,039
CURRENT FILING DATE: 2005-01-11
PRIOR APPLICATION NUMBER: 10/245,871
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 10/197,000
PRIOR FILING DATE: 2002-09-17
PRIOR APPLICATION NUMBER: 09/396,813
PRIOR FILING DATE: 1999-09-14
NUMBER OF SEO ID NOS: 1452
COLUMN APPLICATION NUMBER: 1993-09-14
NUMBER OF SEO ID NOS: 1452
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                                             1 VLHDDLLEA 9
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                                                                                        4; Conservative
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                                                                                        Score 23; DB 7; 1
Pred. No. 1.4e+05;
3; Mismatches 2.
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Pred. No. 1.4e+05;
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RBSULT 15 US-10-485-788A-352 ; Sequence 352, Application US/10485788A

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Publication No. US20050282743A1

GENERAL IMPORMATION:

APPLICANT: Lapeter S.

APPLICANT: Rabinowitz, Joshua D.

APPLICANT: Schweizer, Johannes

APPLICANT: Schweizer, Johannes

APPLICANT: Garrick, Deanna Marie

APPLICANT: Garrick, Deanna Marie

APPLICANT: Carrick, Deanna Marie

CURRENT FILING DATE: 20054-00320US

CURRENT FILING DATE: 20064-02-03

PRIOR APPLICATION NUMBER: US 60/360,061

PRIOR FILING DATE: 2002-02-25

PRIOR FILING DATE: 20
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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seq length: 9
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1: geneseqp1980
2: geneseqp1990
3: geneseqp2000
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Copyright (c) 1993 - 2006
geneseqp2000s:*
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Biocceleration Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

ALIGNMENTS

RESULT 1
AAW99197
ID AAW9
XX AAW9
AC AAW9
XX Mino
XX WO95
XX U71
XX U71
XX WPI
PH Goul
XX WPI
PT dis(
PT mar)
XX WPI
XX Dis(
CC epilo
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XX SQ Seq Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease. Minor histocompatibility antigen HA-1 T-cell epitope #3 23-JUL-1997; 23-JUL-1998; 04-FEB-1999. WO9905174-A1. Homo sapiens. 20-MAY-1999 AAW99197; AAW99197 standard; peptide; 9 AA. (first entry) 97EP-00202303. 98WO-NL000425.

(UYLE-) RIJKSUNIV LEIDEN.

Goulmy EAJM, Hunt DF, Engelhard VH;

WPI; 1999-153312/13.

A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation. HA-1 - useful to treat immune versus graft disease in bone

Disclosure; Page 15; 47pp; English.

The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as a medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the particular it can be used with bone marrow transplantation, in the treatment of severe aplastic anaemia, leukaemia, and immune deficiency

Sequence 9 AA;

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RESULT 3
ADH40334
ID **MADH40334 standard; peptide; 9 AA.
XX
AC ADH46034;
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Best Local S
Matches 9
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Best Local S
Matches 9
                                                                                                                                                                                                              The present sequence represents part of the minor histocompatibility antigen HA-1 R-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of geneetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                                                                                                                                                                                                                                              Typing minor histocompatibility antigen HA-1 - by amplifying and identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. det of genetic aberrances.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW97374 standard;
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                                                                                                                                                                                         Sequence 9
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02-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                         Claim 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                       l Similarity
9; Conserv
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                                                                                                                 VLRDDLLEA
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                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97EP-00202303.
98EP-00870125.
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Pred. No. 2e+06;
; Mismatches 0
                                                                                                                                                    Score 42;
Pred. No.
                                                                                                                                         Mismatches
                                                                                                                                                    DB 2;
2e+06;
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                                                                                                                                                                Length
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                                                                                                                                                                                                                                                                                                                                                                                               detection
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RESULT 4
AAW99196
ID AAW9
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AC AAW9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a novel method for providing epitopes of allelic convariants of antigenic proteins from specific species based on single conclectide polymorphism (SNP), by defining target protein/peptide or its converted to the polymorphism (SNP), by defining target protein/peptide or its converted to the protein dentifying, converted the peptide/protein variants, expression product or its converted to the protein dentifying, converted the protein and protein dentifying, converted the protein binding to MHC protein. A protein of the invention has converted to the invention has converted to the protein and protein 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 9
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minor l
Minor histocompatibility antigen; HA-1; T-cell epitope; immuno graft versus host disease; bone marrow transplant; leukaemia; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 82; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JUN-2002; 2002EP-00013423
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                                                                                                     Minor histocompatibility antigen HA-1 T-cell epitope #2.
                                                                                                                                                                                                                                                    AAW99196 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
9; Conserv
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                                                                                                                                                      (first entry)
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Pred. No. 2e+
0; Mismatches
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                                                          immunological;
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vaccine;

Homo

sapiens

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Matches 8
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WPI; 1999-142960/12
                    Goulmy
                                                               23-JUL-1997;
02-JUN-1998;
                                                                                              23-JUL-1998;
                                                                                                                   04-FEB-1999.
                                                                                                                                         WO9905313-A2
                                                                                                                                                              Homo
                                                                                                                                                                               Intron; minor histocompatibility antigen HA-1; typing allele; H allele; R allele; polymorphic nucleotide; HA-1 typing; bone marrow transplant; severe aplastic anaemia; leukaemia; immune deficiency disease; ss.
                                                                                                                                                                                                                             HA-1 H-allele sequence.
                                                                                                                                                                                                                                                                                            AAW97375 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A new minor histocompatibility antigen, diseases and prevent rejection and host marrow and organ transplantation.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 3;
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                                         (UYLE-) RIJKSUNIV LEIDEN
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                                                                                                                                                             sapiens
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                                                                                                                                                                                                                                                                                                                                                  8
                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                  (first entry)
                                                               97EP-00202303
98EP-00870125
                                                                                              98WO-EP004928
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                                                                                                                                                                                                                                                                                            protein; 9 AA.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
2e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HA-1 -
versus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           useful to treat immune graft disease in bone
                                                                                                                                                                                                                                                                                                                                                                                                                Length 9;
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RESULT 6
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Best Local S
Matches 8
The invention relates to a novel method for providing epitopes of allelic variants of antigenic proteins from specific species based on single nucleotide polymorphism (SNP), by defining target protein, peptide or its subset, screening database of DNA encoding target protein, identifying, selecting allelic peptide/protein variants, expression product or its fragment encoded by DNA sequence having SNP, creating variant epitopes, selecting epitopes binding to MHC protein. A protein of the invention has cytostatic activity, and may have a use in a vaccine. The method is
                                                                                                                                                                                                  Providing allelic variant epitope of protein based on single nucleotide polymorphism by defining target protein, screening database of protein, identifying, selecting allelic variant protein, creating variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents part of the minor histocompatibility antigen HA-1 H-allele. The specification describes methods for typing alleles (preferably the H and R alleles) of the minor histocompatibility antigen HA-1 in a sample, which comprise detecting polymorphic nucleotides in the cDNA or genomic nucleic acids of the alleles. The methods can be used for HA-1 typing for bone marrow transplants, severe aplastic anaemia, leukaemia and immune deficiency diseases, as well as detection of genetic aberrances. The probes and primers of the invention can be used to screen for the HA-1 alleles. The HA-1 peptides can be used anti-idiotypic B cells and/or T cells and antibodies
                                                                                                                                                     Disclosure; Page 82; 119pp; English
                                                                                                                                                                                                                                                                       WPI; 2004-082200/08
                                                                                                                                                                                                                                                                                                      Strittmatter W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human; cytostatic; vaccine; SNP profile; cancer; leukaemia; minor histocomptibility antigen; mHAg; T cell epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Typing minor histocompatibility antigen HA-1 - by amplifying identifying HA-1 R and HA-1 H alleles of HA-1, used for, e.g. of genetic aberrances.
                                                                                                                                                                                                                                                                                                                                                                                                       13-JUN-2003; 2003WO-EP006251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003106692-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human minor histocomptibility antigen HA-1 T cell eptiope.
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                                                                                                                                                                                                                                                                                                                                                                        13-JUN-2002; 2002EP-00013423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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8; Conserv
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88.9%;
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2e+06;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          useful for generating a SNP profile of one or more individuals from a given species by applying the method for several protein from the individuals, where the SNP profile was related to disease, preferably cancer. This is useful for diagnosing a disease in an individual by generating the SNP-related polymorphic profile. A method of the invention is useful for transplanting haematopoletic stem cells from a donor to a recipient and treating cancer, preferably leukaemia, and for determining the progression, regression or onset of a treated disease. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Minor histocompatibility antigen; HA-1; T-cell epitope; immunological; graft versus host disease; bone marrow transplant; leukaemia; vaccine; diagnosis; aplastic anaemia; immune deficiency disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Minor histocompatibility antigen HA-1 T-cell epitope #1
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                              diseases
                                                      The present sequence represents a new peptide (P1) constituting a T-cell epitope obtainable from the minor histocompatibility antigen HA-1. The peptide is immunogenic and can be used as part of a vaccine. P1 is used as medicine, to induce tolerance for transplants, prevent rejection and/or graft versus host disease, or to treat (auto) immune diseases. In particular it can be used with bone marrow transplantation, in the
                                                                                                                                                                                                                                                                                                                                                                                    04-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
Sequence 9 AA;
                                                                                                                                                                                                A new minor histocompatibility antigen, diseases and prevent rejection and host
                                                                                                                                                                                                                                                                                                                              23-JUL-1997;
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                                             treatment of severe
                                                                                                                                                          Claim 1; Page 32; 47pp;
                                                                                                                                                                                                                                            WPI; 1999-153312/13.
                                                                                                                                                                                                                                                                      Goulmy EAJM, Hunt DF,
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8; Conserv
                                                                                                                                                                                     and organ transplantation.
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                                                                                                                                                                                                                                                                                                                              97EP-00202303
                                                                                                                                                                                                                                                                                                                                                          98WO-NL000425
                                                                                                                                                                                                                                                                                                                                                                                                                                          /label= His, Arg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                           aplastic
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                                                                                                                                                          English.
                                                                                                                                                                                                                                                                      Engelhard
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Pred.
                                             anaemia,
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Mismatches
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2e+06;
                                                leukaemia,
                                                                                                                                                                                                   HA-1 - useful to treat immune versus graft disease in bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                and immune
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                                                deficiency
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RANGULT 8
AAW9752
ID AAW9
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Best Local S
Matches 8
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      T-cell epitope; minor histocompatibility antigen HA-1; vaccine; transplant rejection; Graft-versus-Host Disease; autoimmune dis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T-cell epitope from the minor histocompatibility antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-MAY-1999
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                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents an immunogenic peptide constituting a T-cell epitope, obtainable from the minor histocompatibility antigen Ha-1. The peptide can be used in vaccines or pharmaceutical formulations as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neoplastic haematopoietic cell.
                                                                                                                                                                                                                                                         The peptides can also be used to raise antibodies, T-cell receptor, and T-cells
                                                                                                                                                                                                                                                                                                                                                            medicines to induce tolerance for transplants so as to prevent and/or Graft-versus-Host Disease, or to treat autoimmune diseas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-142855/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JUL-1997;
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                                                                                                                                                                                                      Sequence 9
                                                                                                                                                                                                                                                                                                         Neoplastic haematopoietic cells presenting the peptides, in an HLA class I context, can be eliminated after specific recognition of the peptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 39; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     versus-host disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunogenic peptide from minor histocompatibility antigen HA-1 - useful for inducing tolerance to transplants and prevent rejection or graft-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9905173-A1
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8; Conserv
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VLXDDLLEA
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88.9%;
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Pred. No. 2e+06;
0; Mismatches
                                                                                                     0;
                                                                                                                                Score 36;
Pred. No.
                                                                                                     Mismatches
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                                                                                                                              2e+06;
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                                                                                                                                                                                                                                                                                                                                                                                          rejection
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RESULT 9
ADT73023
ID ADT7

ADT73023 standard; peptide; 9 AA.

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RESULT 10
AAW47427
ID AAW47
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AC AAW47
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DT 05-JU
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                                                                                                                                                                                                                                                                                                                                              This invention relates to novel isolated or purified peptides of the CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins: non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein I and CC SH (also known as the 1A proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC composition comprising an above peptide mixed with a pharmaceutical CC excipient or an RSV immunogenic composition comprising a recombinant CC expression vector with a nucleic acid insert encoding an above peptide. CC Thymphocytes (CTIs) that respond to a major histocompatibility complex CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC RSV peptide is useful for preparing a diagnostic composition or an RSV (prophylactic or therapeutic) vaccine composition for a DNA-based CC immunisation, or for preparing an immune response provoking vaccine in CC the event of RSV infection (the vaccine being prepared by contacting the CC polypeptide in an immune response-provoking anount of specific CTI).

CC Accordingly, these peptide compositions have virucidal activity. This cc invention.
                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 4
 Prenyl diphosphate synthetase preserved region VI sequence.
                                 05-JUN-1998
                                                                 AAW47427;
                                                                                                                                                                                                                                                                                                          Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
                                                                                               AAW47427 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 16; SEQ ID NO 919; 143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lasters I, Desmet J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-APR-2003; 2003EP-00447095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-APR-2004; 2004WO-EP004061.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human respiratory syncytial virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human RSV L high affinity binding peptide SeqID 919
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human respiratory syncytial virus; cytotoxic T lymphocyte; CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            epitope;
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                                                                                                                                                                                                            1 VLRDDLLEA 9
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                               (first entry)
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                                                                                               peptide; 9
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                                                                                                                                                                                                                                                         Score 29;
Pred. No.
                                                                                                                                                                                                                                            Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RSV infection; virucidal; vaccine
                                                                                                                                                                                                                                                           2e+06;
                                                                                                                                                                                                                                                                          DB 8;
                                                                                                                                                                                                                                                                        Length 9;
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RESULT 11
ADT73690
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence from preserved region VI of prenyl diphosphate synthetase (PDS), was used in the preparation of primers for the amplification of PDS DNA. Substances synthesised by PDS, i.e. polyprenyl diphosphates, are precursors of physiologically active substances, e.g. vitamin K and ubiquinones
                  WPI; 2004-758334/74.
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA encoding prenyl diphosphate synthetase subunit(s) - new Micrococco prenyl diphosphate synthetase subunit polypeptide(s), and methods for
                                                                                                   16-APR-2004; 2004WO-EP004061.
                                                                                                                                            WO2004092207-A2
                                                                                                                                                                Human respiratory syncytial virus
                                                                                                                                                                                  human respiratory syncytial virus; cytotoxic T lymphocyte;
MHC epitope; DNA-based immunisation; RSV infection; virucio
                                                                                                                                                                                                                   Human
                                                                                                                                                                                                                                        13-JAN-2005
                                                                                                                                                                                                                                                           ADT73690;
                                                                                                                                                                                                                                                                               ADT73690 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Muramatsu M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prenyl diphosphate synthetase; polyprenyl diphosphate; vitamin ubiquinone; preserved region VI.
                                       Lasters
                                                                               16-APR-2003; 2003EP-00447095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        preparing enzymes from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUN-1997;
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                                                          (ALGO-) ALGONOMICS
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                                                                                                                                                                                                                 RSV L high affinity binding peptide SeqID 1586
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                                     Desmet J,
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                                                                                                                                                                                                                                                                               peptide; 9
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57.1%;
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                                      Stegmann
                                                                                                                                                                                   immunisation; RSV infection; virucidal; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           subunit(s).
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Pred. No.
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New respiratory syncytial virus (RSV) peptides (e.g.

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M2-1, M2-2,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 4
                             Disclosure; SEQ ID NO 663; 143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADT72767 standard; peptide; 9
                                                      New respiratory syncytial virus (RSV) peptides (e.g. G, M, M2-1, M2-2, or P) for inducing an immune response to RSV or for diagnosing, preventing or treating viral infections, particularly RSV infection.
                                                                                                                                                                                                                                                                                                  WO2004092207-A2
                                                                                                                                                                                                                                                                                                                                                                                                       Human RSV L high affinity binding peptide SeqID 663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADT72767;
                                                                                                                                                                                                                                       16-APR-2004; 2004WO-EP004061
                                                                                                                                                                                                                                                                                                                            Human respiratory syncytial virus
                                                                                                                                                                                                                                                                                                                                                           MHC epitope; DNA-based immunisation;
                                                                                                                                                                                                                                                                                                                                                                           human respiratory
                                                                                                                    WPI; 2004-758334/74
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                                                                                                                                                                                                                                                                                                                                                             syncytial virus; cytotoxic T lymphocyte; CTL;
based immunisation; RSV infection; virucidal; vaccine
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This invention relates to novel isolated or purified peptides of the

Sequences AAU71093-AAU71796 represent the 103P3E8-related protein peptide fragments of the protein. 103P3E8 exhibits tissue specific expression in normal adult tissue, but it is also aberrantly expression.

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e.g.

Disclosure; Page 90; 128pp; English.

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CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein L and CC eight structural proteins preferably G, also M, M2-1, M2-2, N, p, p, and CC composition comprising an above peptide mixed with a pharmaceutical CC excipient or an RSV immunogenic composition comprising an above peptide cC expression vector with a nucleic acid insert encoding an above peptide. CC The present invention describes an in vitro method of detecting cytotoxic CC Thymphocytes (CTLs) that respond to a major histocompatibility complex CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The CC (Prophylactic or therapeutic) vaccine composition for a DNA-based CC (Prophylactic or therapeutic) control (the vaccine being prepared by contacting the vent of RSV infection (the vaccine being prepared by contacting the CC polypeptide in an immune response provoking amount of specific CTL).

CC Accordingly, these peptide compositions have virucidal activity. This competition is a human RSV high binding affinity peptide of the
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Best Local Similarity
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                                                                                                              Monitoring 103P3E8 gene products in sample from patient (suspected of) having cancer, useful for diagnosing, managing or treating cancers, e. prostate cancer, comprises determining presence of aberrant 103P3E8 ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      103P3E8; prostate; bladder; kidney; colon; lung; breast; rectum; stomach; tumour; cancer; cytostatic; gene therapy; antibody therapy; ribozyme; single chain monoclonal antibody; serum; blood; urine; tissue; human;
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                                                                                                                                                                                  WPI; 2002-061976/08.
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Pred. No.
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2e+06;
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RESULT 14
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encoding II separate viral proteins: non-structural proteins NS-1 (also known as the IC protein) & NS-2 (IB protein), a polymerase protein I and eight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and SH (also known as the IA protein). Specifically, it refers to a composition comprising an above peptide mixed with a pharmaceutical excipient or an RSV immunogenic composition comprising a recombinant expression vector with a nucleic acid insert encoding an above peptide. The present invention describes an in vitro method of detecting cycotoxic T lymphocytes (CTLs) that respond to a major histocompatibility complex (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The RSV peptide is useful for preparing a diagnostic composition or an RSV (prophylactic or therapeutic) vaccine composition for a DNA-based
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            many cancers including tumours of the prostate, bladder, kidney, colon, lung, breast, rectum and stomach. The 103938 polymucleotide, its relate protein and peptide fragments and specific PCR primers are therefore useful for diagnosing and treating cancer. A vector comprising a polymucleotide which encodes a single chain monoclonal antibody, that immunospecifically binds to an 1039388 related protein, and a ribozyme capable of cleaving a polymucleotide having the 1039388 coding sequence, are both useful in the preparation of a composition for treating a patient with a cancer that expresses 1039388. The sequences can be used in diagnostic methods to monitor the level of 1039388 gene products in secun, blood, urine and tissue and to thereby detect the presence of
                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to novel isolated or purified peptides of the human respiratory symcytial virus (RSV), in particular ten RSV genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       preventing
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RESULT 15
ADT73689
ID ADT73
CC This invention relates to novel isolated or purified peptides of the CC human respiratory syncytial virus (RSV), in particular ten RSV genes CC encoding 11 separate viral proteins: non-structural proteins NS-1 (also CC known as the 1C protein) & NS-2 (1B protein), a polymerase protein L and CC sight structural proteins preferably G, also M, M2-1, M2-2, N, P, F, and CC SH (also known as the 1A protein). Specifically, it refers to a CC composition comprising an above peptide mixed with a pharmaceutical CC expression vector with a nucleic acid insert encoding an above peptide. CC The present invention describes an in vitro method of detecting cytotoxic CC Tlymphocytes (CTLs) that respond to a major histocompatibility complex CC (MHC) class I restricted T-cell stimulating peptide epitope of RSV. The RSV peptide is useful for preparing a diagnostic composition or an RSV CC (prophylactic or therapeutic) vaccine composition for a DNA-based CC immunisation, or for preparing an immune response provoking vaccine in CC the event of RSV infection (the vaccine being prepared by contacting the CC polypeptide in an immune response-provoking amount of specific CTL). Accordingly, these peptide compositions have virucidal activity. This contacting is a human RSV high binding affinity peptide of the
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                                                                                                                                                                                                                                                                                                                                                                                          Claim 16; SEQ ID NO 1585; 143pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human RSV L high affinity binding peptide SeqID 1585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lasters I, Desmet J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-APR-2003; 2003EP-00447095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-APR-2004; 2004WO-EP004061
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human respiratory syncytial virus; cytotoxic T lymphocyte; CTL;
MHC epitope; DNA-based immunisation; RSV infection; virucidal; vaccine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADT73689;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADT73689 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunisation, or for preparing an immune response provoking vaccine in the event of RSV infection (the vaccine being prepared by contacting the polypeptide in an immune response-provoking amount of specific CTL). Accordingly, these peptide compositions have virucidal activity. This peptide sequence is a human RSV high binding affinity peptide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9
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42.9%;
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Pred. No.
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Query Match Semilarity 42.9%; Excre 25; DB 6; Length 9;
Ber Locil Semilarity 42.9%; Pred. No. 22+06;
Matches 3; Conservative 4; Mismatches 0; Indels 0; Gape 0;
Db 2:INDDL 9
Search completed April 6, 2006, 16:19:10
Job time: 112.667 secs
```

GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.

OM protein - protein search, using sw model

Run on: April 6, 2006, 16:19:36; Search time 18.6667 Seconds (without alignments) 46.390 Million cell updates/sec

Title: Perfect score: Sequence: US-10-791-217A-5 42 1 VLRDDLLEA 9

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters:

791

Minimum DB seq length: 0 Maximum DB seq length: 9

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : PIR 80:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

222 28 29	18 22 22 24	1 2 3 3 3 3 5 5 6 6 7 7 7 11 11 11 11 11 11 11 11 11 11 11	
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T11779 S25266 PT0602 S22428 B33099	A61328 PH0108 PT0562 B30572 B30572 A39841 A37027 D60274	S68004 PC4131 PC4131 PC4134 S20446 A60427 A60427 PT0669 PT0669 PT0669 PT0533 PT0533 PT0533 PT0533 PT0533 PT0533 PT0722 PT0722 PT0722	
phosphoglycerat pilE protein - T-cell receptor chitin-binding	profin	hucolin, 75k hucolin kinase elastase per elastase cytopic forcell receptor reell receptor reell receptor acetylcholines calsequestrin, 7-cell receptor reell receptor rec	Description
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H48394 S78024	A26209 A35779 B44510	A44787 A48360	B57444 C57444	PS0253 A12872	PW0002 B39841	B28495	869165
glycoprotein compo ribosomal protein	protein-glutamine neuropeptide Antho hypothetical prote	callifMRFamide 10 gamma subunit of P	neuropeptide Grb-A neuropeptide Grb-A	glycine cleavage s transaldolase (EC	chlorophyll a/b-bi dextransucrase (EC	conopressin S - co	ferredoxin a2 - Ja

ALIGNMENTS

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RESULT 4
S20446
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C;Species: Rattus norvegicus
C;Date: 25-Aug-2003 #sequence_revision 25-Aug-2003 #text_change 25-Aug-2003
C;Accession: A59489
                                                                                                                                                                                    macrophage cytotoxicity-inducing factor, 29K - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 12-Feb-1993 #sequence_revision 12-Feb-1993 #text_change 09-Jul-2004
C;Accession: A60427
R;Jones, C.M.; Prince, C.A.; Williams, J.S.
Exp. Hematol. 19, 704-709, 1991
A;Title: Purification and amino acid analysis of a human macrophage cytotoxicity-inducin A;Reference number: A60427; MUID:91372335; PMID:1909970
A;Accession: A60427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              elastase - Pseudomonas aeruginosa (;Species: Pseudomonas aeruginosa (;Species: Pseudomonas aeruginosa (;Date: 19-Mar-1997 #sequence_revision 24-Jul-1997 #text_change 24-Jul-1997 (;Accession: S20446 R;Kessler, E.; Safrin, M.; Peretz, M.; Burstein, Y. FEBS Lett. 299, 291-293, 1992 A;Title: Identification of cleavage sites involved in proteolytic processing of Pseudomo A;Reference number: S20446; MUID:92183956; PMID:1544509 A;Accession: S20446
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                                                                                           A;Cross-references: UNIPROT:Q7M4R5; UNIPARC:UPI0000142BE0
A;Note: the sequence from the text on page 706 is inconsistent with that from page C;Keywords: cytokine
                                                                                                                                                 A; Molecule type: proto
A; Residues: 1-9 < JON>
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A;Cross-references:
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                              33.3%;
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              Score 14; DB 2;
Pred. No. 2.8e+05;
0; Mismatches 1
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Pred. No.
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Pred. No. 2.8e+05;
2; Mismatches 0
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C;Date: 09-Oct-1992 #sequence revision 09-Oct-1992
C;Accession: PH0942
R;Gold, D.P.; Offner, H.; Sun, D.; Willey, S.; Vande
J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta chains in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 6
PH0942
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C;Species: Citrobacter freundii
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 09-Jul-2004
C;Accession: I40697
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Experimental source: complete Freund's A;Note: the authors translated the codon C;Keywords: T-cell receptor
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A; Residues: 1-9 <GOL>
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                                                                                                                                                                                            C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0679; PT0708
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Shiuan, D.; Campbell, A.
Gene 67, 203-211, 1988
A;Title: Transcriptional regulation and gene arrangement
A;Reference number: I40697; MUID:89006280; PMID:2971595
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                                                                                                               R;Feeney, A.J.
1. Exp. Med. 174, 115-124, 1991
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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                                    A; Molecule type: DNA
A; Residues: 1-5 <FEE>
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  A;Experimental source:
                  A; Cross-references: UNIPARC: UPI000017C84D
                                                                          A; Status: translation not
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llarity 100.0%;
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day 18 fetal thymus,
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                                                                               shown
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Pred. No.
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3. 2.8e+05;
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. 2.8e+05;
strain BALB/c, 154-2J
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T-cell receptor beta chain V-D-J region (clone 13) - rat (fragment)
                                                                                     adjuvant-immunized lymph TGC for residue 2 as Ala
                                                                                                                                                                                                                                                      S.; Vandenbark, A.A.; Wilson, D.B.
                                                                                                                                                                                                                                                                                          #text_change 30-May-1997
                                  Length 9;
Indels
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R; Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A; Title: Junctional sequences of fetal T cell receptor beta A; Reference number: PT0509; MUID:91277601; PMID:1711558
A; Accession: PT0601
A; Accession: PT0601
                                                                        Cerebellar degeneration-related protein - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 28-Sep-1990 #sequence_revision 28-Sep-1990 #text_change 24-Jun-1993
C;Accession: B35640
R;Chen, Y.T.; Rettig, W.J.; Yenamandra, A.K.; Kozak, C.A.; Chaganti, R.S.K.; Posner, Proc. Natl. Acad. Sci. U.S.A. 87, 3077-3081, 1990
A;Title: Cerebellar degeneration-related antigen: a highly conserved neuroectodermal A;Reference number: A35640; MUID:90222173; PMID:2326268
A;Accession: B35640
                                                                                                                                                                                                                                                            RESULT 10
B35640
                    A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-6 <CHE>
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 09-Jul-2004
C;Accession: PT0601; PT0617; PT0694
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A;Experimental source: newborn thymus, strain BALB/c, 161-2B
C;Keywords: T-cell receptor
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A;Status: translation not shown
A;Molecule type: DNA
A;Residues: 1-5 <PE2>
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C;Keywords: T-cell receptor
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A;Cross-references: UNIPARC:UPI000017C63A
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A;Experimental source: day 18 fetal thymus,
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A;Experimental source: newborn thymus,
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A; Residues: 1-5 < FEE>
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;Residues: 1-5 <FE3>
;Cross-references: UNIPARC:UPI000017C7E6
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Best Local
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Pred. No. 2.8e+05;
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Query Match

Score 12;

DB 2;

Length 6;

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R;Cala, S.E.; Jones, L.R.
Biol. Chem. 266, 391-398, 1.
A;Title: Phosphorylation of ca.
A;Reference number: A39040; MU
A;Accession: B39040
A;Status: preliminary
                                                                                                              RESULT 13
B39040
calsequestrin, fast skeletal muscle - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 23-Aug-1991 #sequence_revision 23-Aug-1991 #tex
C;Accession: B39040
C;Accession: B39040
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J. Biol. Chem. 263, 1140-1145, 1988
A;Title: Divergence in primary structure between the molecular
A;Reference number: A34026; MUID:88087239; PMID:3335534
A;Accession: A34026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acetylcholinesterase (EC 3.1.1.7) 5.6S form - Pacific electric ray (fragment) C;Species: Torpedo californica (Pacific electric ray) C;Date: 16-Mar-1990 #sequence_revision 16-Mar-1990 #text_change 08-Nov-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Keywords: alternative splicing; carboxylic ester hydrolase
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A;Experimental source: adult thymus, strain BALB/c
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A; Residues: 1-6 < FEE>
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C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
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                                         cardiac and skeletal muscle MUID:91093153; PMID:1985907
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Pred. No. 2.8e+05;
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Pred. No.
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1; Mismatches 0
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R; Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A; Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A; Reference number: PT0509; MUID:91277601; PMID:1711558
A; Accession: PT0628
A; Status: translation not shown
A; Molecule type: mRNA
A; Residues: 1-7 <FEE>
A; Cross-references: UNIPARC:UPI000017C7CD
A; Experimental source: newborn thymus, strain BALB/c
C; Keywords: T-cell receptor
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PT0628
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PT0722
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0628
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A;Cross-references: UNIPARC:UPI000017C5BC
C;Keywords: phosphoprotein; skeletal muscle
                                                                                                                                                                                                                                                                                                             C;Accession: PT0722

R;Feeney, A.J.

J. Exp. Med. 174, 115-124, 1991

J.;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.

A;Reference number: PT0509; MUID:91277601; PMID:1711558

A;Accession: PT0722

A;Status: translation not shown

A;Residues: 1-7 <FEE>

A;Residues: 1-7 <FEE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T-cell receptor beta chain V-D-J region (135-1G) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0722
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A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
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Search completed: April 6, 2006, 16:26:10 Job time : 18.6667 secs
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Matches
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Q70SM2_HUMAN
Q7KYP6_HUMAN
Q8LPT5_MAIZE
P82568_STRPY
Q85710 9REFR
Q80TD7 9HIV1
Q15897 HUMAN
Q9HDS4 ASPPL
Q5RLS1 PIG
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Q93CR0 STAEP
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 MEDLINE=22779048; PubMed=12897253; DOI=10.1105/tpc.012526;
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O1-FEB-2005 (Tr Phytoene syntha Name-psy2; Zea mays (Maize Eukaryota; Viri. Spermatophyta; I Spermatophyta; I NCBI TaxID=4577 [1] NUCLEOTIDE SEQU STRAIN=Y-13, an MEDLINE=2777904	TT 2 P7 MAIZ Q7X8P7 Q7X8P7 01-OCT		ry Mat t Loca ches	NUCLEOTII STRAIN=Y MEDLINE=Y Palaisa I "Contrass disequil Plant Ce. EMBL; AY NON TER NON TER SEQUENCE	01-OCT 01-OCT 01-OCT 01-OCT Phytoe Name=p Zea ma Eukary Sperma Sperma PACCAD NCBI_T	ILT 1 103 MAIZE Q7XB03 MAIZE Q7XB03;	
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2005 (TrKMBLTel. 29,) e Synthase 2 (Fragmeni 12); s (Maize). ta; Viridiplantae; St. ta; Viridiplantae; St. ta; Magnoliophyta clade; Panicoideae; Au clade; Panicoideae; Au xID=4577; IDB SEQUENCE. y-13, and W-16; =22779048. PubMed=128	MAIZE PRELIMINARY; 2003 (TremBLrel. 2	RD 4	ilarity Conservat	SEQUENCE. 79048; Pub 79048; Pub 9: effects 10: 1795-18 612; AAP55 1 8 AA; 915	01-OCT-2003 (TREMBLIRE1. 25, Creat 01-OCT-2003 (TREMBLIRE1. 25, Last 01-OCT-2003 (TREMBLIRE1. 25, Last 01-OCT-2003 (TREMBLIRE1. 25, Last Phytoene synthase 2 (Fragment). Name=psy2; Zea mays (Maize). Zea mays (Maize). Eukaryota; Viridiplantae; Strepto Spermatophyta; Magnooliophyta; Li. PACCAD olade; Panicoideae; Androy NCBI_TaxID=4577;	B PRELIMII	31.0 31.0 31.0 31.0 31.0 31.0 31.0 31.0
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men men st st yta ;; A	ິທິທີ		* *	128 12003 17006	ymer yta		00000000000000
L). c). c). c). c). c). c). c).	Y; PRT; 9 AA. 25, Created) 25, Last sequence update)		Score 18; DB 2; Length 8 Pred. No. 2.2e+06; 1; Mismatches 0; Indel); DOI=10.1105/tpc.01 ams M., Rafalski A.; i on sequence diversi /nthase loci."; enomic_DNA.	, Created) , Last sequence update) , Last annotation update) ent). Streptophyta; Embryophyta; T. Striptopsida; Poales; Poace hadropogoneae; Zea.	ALIGNMENTS PRT; 8 AA.	Q6PUD7_SV40 Q6PUD9_SV40 Q6PUD11_SV40 Q6PUD11_SV40 Q6QSX9_SV40 Q6QSX9_SV40 Q6QA1X4_FUNHE IPYR_RHOVI Q7RY8_PHOVI Q7RY8_SHOVI Q6UVK2_MALDO Q56ST9_SAWCA Q56ST1_9DIPS Q56ST1_9DIPS Q56ST1_9DIPS Q56ST1_YIBOP Q56ST3_VIBOP
; Tracheophyta; oaceae;			18; lels 0; Gaps 0,	.2526; .ty and linkage	Tracheophyta;		Q6pud7 simian viru Q6pud9 simian viru Q6pud9 simian viru Q6qsy9 simian viru Q64ix4 fundulus he P82992 rhodopseudo Q7ry89 neurospora Q61cv2 homo sapien Q6uvk2 malus domes Q56s89 sambucus ca Q56st1 viburnum ur Q56st2 viburnum tr Q56st2 viburnum cp Q56st3 viburnum op Q56st3 viburnum op

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RESULT 3
Q705M2 HUMAN PRELIMINARY;
AC Q705M2;
DT 05-UUL-2004 (TrEMBLrel. 27,
DE Hypermethylated in cancer 1
GN Name=HIC1;
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordat
OC Mammalia; Eutheria; Euarcho
OC Homo.
OX NCBI_TaxID=9606;
RN [1]
RP NUCLEOTIDE SEQUENCE.
RC TISSUE=Mammary gland;
RA PINCLEOTIDE SEQUENCE.
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RT COCOGEON 23:4023-4031(2004)
DR EMBL; AJ550616; CAD79467.1;
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SQ SEQUENCE 9 AA; 964 MW;
Q7KYP6 HUMAN
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pinte S., Guerardel C., Deltour S., Godwin A.K.,
pinte S., Guerardel C., Deltour S., Godwin A.K.,
"Identification of a second G-C-rich promoter cor
murine and rat tumor suppressor genes HIC1.";
Oncogene 23:4023-4031(2004).
EMBL; AJ550616; CAD79467.1; -; mRNA.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini; Homin
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EMBL; AY300665; AAP55403.1; -; Genomic_DNA.
NON_TER 1
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                             TISSUE=Miltenberger class V;
MEDLINE=90005483; PubMed=2792104;
Vignal A., Rahuel C., El-Maliki B
                                                                                                                                                                                          05-UIL-2004 (TREMBLrel. 27, Created)
05-UIL-2004 (TREMBLrel. 27, Last sequence update)
05-UIL-2004 (TREMBLrel. 27, Last annotation updat
Mi.V glycoprotein (exon P3B) (Fragment).
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                              NUCLEOTIDE SEQUENCE.
                                                                                                             NCBI_TaxID=9606;
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                                                                                                                                             Chordata; Craniata; Vertebrata; Euteleostomi; Euarchontoglires; Primates; Catarrhini; Hominidae;
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1; Mismatches 0
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RESULT 5
QBLPTS MAIZE
QBLPTS MAIZE
ID QBLPTS;
AC QBLPTS;
AC QBLPTS;
DT 01-OCT-2002 (TIEMBLIEL 22,
DT 01-OCT-2002 (TIEMBLIEL)
CE BACADOLOGIA (TIEMBLIEL)
CE BACADOLOGIA (Maize)
CE GRACAD Clade; Panicoideae; SOC Spermatophyta; Magnoliophyt
CC PACCAD Clade; Panicoideae;
CX NCBL TAXID=4577;
RN NUCLEOTIDE SEQUENCE.
RA Ching A.S., Caldwell K.S.,
RA MOTGGATE M., RAFALSKI J.A.;
RA MOTGGATE M., RAFALSKI J.A.;
RA EMBL; AY094310; AAM21836.1;
FT NON TER 1
SQ SEQUENCE 9 AA; 977 MW;
RESULT 6
P82568 STRPY
1D P82568;
AC P82568;
DT 01-0CT-2000 (T
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DE Unknown protect
OS Streptococcus
OC Bacteria; Firm
OC Streptococcus
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RN [1]
RP PROTEIN SEQUEN
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Eur. J. Biochem. 184:337-344(1989)
[2]
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01-OCT-2002 (TrEMBLrel. 22,
Beta-expansin-like protein can mays (Maize).
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Vignal A., Rahuel C., El-Maliki
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Ching A.S., Caldwell K.S., Jung M.,
Morgante M., Rafalski J.A.;
Submitted (APR-2002) to the EMBL/Ger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bukaryota; Viridiplantae; Streptophyta; En
Spermatophyta; Magnoliophyta; Liliopsida;
PACCAD clade; Panicoideae; Andropogoneae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL;
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                                                                                                                                                01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence up
01-OCT-2003 (TrEMBLrel. 25, Last annotation
Unknown protein from 2D-page (Fragment).
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                                                                         Streptococcus.
                                                                                                      Bacteria; Firmicutes;
                                                                                                                           Streptococcus pyogenes.
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                                               NCBI_TaxID=1314;
  PROTEIN SEQUENCE,
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3; Conserv
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     AND MASS SPECTROMETRY
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33.3%;
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                                                                                                         Lactobacillales; Streptococcaceae;
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(Fragment).
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Pred. No.
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Pred. No. 2.2e+06;
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Best Local S
Matches 2
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Best Local
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Viruses; ssRNA negative-strand viruses; Paramyxoviridae; Paramyxovirinae; Avulav NCBI_TaxID=11176;
                              Newcastle disease virus.
                                       Name=HN/HN0
                                                  ONH/NH
                                                         01-JUN-2003 (TrEMBLrel.
01-JUN-2003 (TrEMBLrel.
                                                                             Q80H91;
01-JUN-2003
                                                                                           QBOH91_9PARA PRELIMINARY;
                                                                                                                                                                                                                                                                  Submitted (JUL-1994) to the EMBL/GenBank/DDBJ databases. EMBL; X80041; CAA56350.1; -; Genomic_DNA.
                                                                                                                                                                                                                                                                                                                           MEDLINE=95255652; PubMed=7737502; DOI=10.1016/0378-1119(94)00530-6; Asturias J.A., Diaz E., Timmis K.N.; "Evolutionary relationship of the biphenyl dioxygenase of the grampositve bacterium Rhodococcus globerulus P6 to multicomponent dioxygenases of gram-negative bacteria.";
                                                                                                                                                                                                                            SEQUENCE
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01-FEB-1997 (TrEMBLrel. 02, Last sequence up
01-JUN-2003 (TrEMBLrel. 24, Last annotation
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Submitted (MAY-2000) to
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NCBI_TaxID=33008;
                                                                                                                                                                                                                                                                                                                                                                                                                 Rhodococcus globerulus.
Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
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MW=22592.04;
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Last annotation updat
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Pred. No. 2.2e
4; Mismatches
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Pred. No.
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RESULT 10
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MUCLEOTIDE SEQUENCE.
MEDLINE=94175786; PubMed=8129624;
Collins M.S., Strong I., Alexander D.J.;
"Evaluation of the molecular basis of parameters of parameters of parameters of parameters of the molecular basis of the molecul
                                                                                                                    Bacteriophage phi-10.
Viruses; dsRNA viruses;
                                                                                                                                                                                                               01-NOV-1999
01-DEC-2001
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Q9XJNO;
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EMBL; Y10620; CAA71631.1; -; Genomic DNA.
GO; GO:0016789; F:hydrolase activity; NAS.
SEQUENCE 9 AA; 1258 MW; C52D07340AB41412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUCLEOTIDE SEQUENCE.
MEDLINE=97336048; PubMed=9192843; DOI=10.1006/geno.1997.4731;
MEDLINE=97336048; PubMed=9192843; DOI=10.1006/geno.1997.4731;
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EMBL; 869419; AAP19628.1; -; Genomic_DNA.
NON_TER 1
NUCLEOTIDE SEQUENCE
MEDLINE=99350412; Po
                                                                                                                                                                                      P10 (Fragment)
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Kere J., Ropers H.H., Kalscheuer V.M.;
"Monoallelic expression of human PEG1/MEST is
specific methylation in fetuses.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human
Eukaryota; Metazoa;
Mammalia; Eutheria;
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Q9UE26;
                                                                                 NCBI_TaxID=90889;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Name=PEG1/MEST;
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2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VRRDRL
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19,
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Pred. No.
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"Isolation of additional double-stranded RNA.";
                                                                                                                                       Eukaryota; Viridiplantae; Strepto
Jungermanniopsida; Metzgeriidae;
Pallaviciniaceae; Jensenia.
MCBI_TaxID=280534;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Forrest L.L., Schuette S.W., Crandall-Stotler "A Molecular Study of the Simple Thalloid Live (Marchantiophyta, Pallaviciniaceae)."; Bryologist 108:204-211(2005).
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Forrest L.L., Schuette S.W., Crandall-Stotler "A Molecular Study of the Simple Thalloid Live (Marchantiophyta, Pallaviciniaceae).";
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79 9MARC
Q4QWV9 9MARC PRELIMINARY;
Q4QWV9;
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Pallavicinia rubristipa
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Forrest L.L., Schuette
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Jungermanniopsida; Metzgeriidae; Metzgeriales; Pallaviciniineae;
Pallaviciniaceae; Pallavicinia.
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BL; AF125675; AAD22555.1; -; Genomic_RNA.
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MEDLINE-86223772; PubMed-2423502; Weber P.C., Palchaudhuri S.; "Incompatibility repressor in a Repplasmid ColV2-K94."; J. Bacteriol. 166:1106-1112(1986). EMBL; M13472; AAA23194.1; -; Genomi Plasmid.
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EMBL; AY734726; AAX46211.1; -
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Drews O., Weiss W., Reil G., Parlar H., Wait R., Goerg A.;
"High pressure effects steep-wise altered protein expression in
Lactobacillus sanfranciscensis.";
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STRAIN=DSM 20451;
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01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
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01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Unknown protein from 2D-page (Fragment).
Lactobacillus sanfranciscensis (Lactobacillus sanfrancisco).
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NCBI_TaxID=2458;
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RESULT 15

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ID Q6KPX4;
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AC Q8KPX4;
DT 01-OCT-2002 (TrEMBLrel. 22, Created)
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DT 01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
DT 01-OCT-2002 (TrEMBLrel. 26, Last sequence update)
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DR Microcypatis sp. T96-1.
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NUCLEOTIDE SEQUENCE.

MEDLINE=22338268; PubMed=12450830;

DOI=10.1128/ARM.68.12.6070-6075.2002;

Baker J.A., Entsch B., Neilan B.A., McKay D.B.;

Baker J.A., Entsch B., Neilan B.A., McKay D.B.;

"Monitoring changing toxigenicity of a cyanobacterial bloom molecular methods.";

Appl. Environ. Microbiol. 68:6070-6076(2002).

EMBL; AY117046; AAM54719.1; -; Genomic_DNA.

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SEQUENCE 8 AA; 890 MW; F4DB01A73771A336 CRC64;
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        length: 0 length: 9
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Match Length
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1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
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first 45 summaries
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US-09-489-760-5
US-09-489-760-2
US-09-489-760-2
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US-09-269-250E-29
US-09-217-609A-11
US-08-873-235B-11
US-08-873-235B-11
US-08-873-235B-11
US-08-89-760-4
US-08-489-760-4
US-08-489-760-4
US-08-399-411-75
US-08-516-859A-20
US-08-516-859A-75
US-08-516-859A-75
US-08-516-859A-75
US-08-516-859A-75
US-08-516-859A-75
US-08-516-859A-75
US-08-516-859A-75
US-08-516-472-75
US-08-459-568-72
US-08-399-411-72
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Sequence 21, Appli
Sequence 21, Appli
Sequence 11, Appli
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Sequence 14, Appli
Sequence 20, Appli
Sequence 70, Appli
Sequence 75, Appli
Sequence 77, Appli
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Best Local :
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  TYPE: PRT ORGANISM:
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TYPE: PRT
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72, 78, 78, 72, 78, 78, 78, 78, 78, 78, 78, 78, 78, 78	App	Appl	App1	App	App	App:	Αp	Αp	Αp	Αp	Ąpp	App	App	App	App	App	App

ALIGNMENTS

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GENERAL INFORMATION:

APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
ITITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT FILING DATE: 2000-01-21
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER APPLICATION UNUBER: PCT/NL98/00424
EARLIER APPLICATION UNUBER: DCT/NL98/00424
EARLIER APPLICATION UNUBER: DCT/NL98/00424
EARLIER APPLICATION UNUBER: DCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-489-760-5
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Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: Goulimy, Elsa
TITLE OF INVERTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN |
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 18
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 5, Application US/09489760 Patent No. 6878375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
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KIAA0223 partial complementary DNA
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Conservative 0
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Pred. No. 4.6e+05;
0; Mismatches 0;
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                                                                                                                                                                     ; ORGANISM: Histocompatibility antigen US-09-489-760-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20, Application US/09269250E
PATENT NO. 6830883
GENERAL INFORMATION:
APPLICANT: GOLLINY, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/09489760
Patent No. 6878375
GENERAL INFORMATION:
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Matches 9; Conserv
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SEQ ID NO 2
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Best Local
                                                                                                          Matches
                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
                                                                                                                                                                                                                                                                         FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
EARLIER FILING DATE: 1998-07-23
                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 16
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Pred. No. 4.6e+05;
0; Mismatches 1
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
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GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Hard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
CURRENT APPLICATION NUMBER: US/09/489,760
CURRENT FILING DATE: 2000-01-21
EARLIER APPLICATION NUMBER: PCT/NL98/00424
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RESULT 7
US-09-217-609A-11
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Patent No. 6830883
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 199-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 29
LENGTH: 9
TYPE: PRT
ORGANIEM: Human
                                                                                                                                 Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                EARLIER FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/09489760 Patent No. 6878375
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa represents a histidine
                                                                                                                                                                                                                  NAME/KEY: PEPTIDE
LOCATION: (3)
OTHER INFORMATION: AMINO ACID X REPRESENTS A HISTIDINE OR AN ARGININE
OTHER INFORMATION: RESIDUE
                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Histocompatibility antigen FEATURE:
                                                                                                                                                                                                                                                                                                                                   LENGTH: 9
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8; Conserv
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ilarity 88.9%;
Conservative
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88.9%;
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Pred. No. 4.6e+05;
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Sequence 11, Application US/09217609A Patent No. 6071733
GENERAL INFORMATION:

APPLICANT:

MURAMATSU, Masayoshi

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                                                                                                                                                                                                                                                                                                                                                                               US-08-873-235B-11
                                                                                                                                                                                                                                                                                                                                          Sequence 11,
Patent No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 64.3%; Score 27; DB 2; Best Local Similarity 57.1%; Pred. No. 4.6e+05; Matches 4; Conservative 3; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/873,235
FILING DATE: 11-Jun-1997
ATTORNEY/AGENT INFORMATION:
NAME: TOFFENETTI, Judith L.
REGISTRATION NUMBER: 39,048
REFERENCE/DOCKET NUMBER: 10235/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-429-176
                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: 202-429-1776
TELEFAX: 202-429-0796
INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                     APPLICANT: SHIMIZU, Naoto
APPLICANT: CHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
CORRESSPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS 6.2
SOFTWARE: WordPerfect 6.1 for Windov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: CHO, Yenwin
TITLE OF INVENTION: Prenyl Diphosphate Synthetase Genes
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
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                                                          COUNTRY: U
ZIP: 20036
                                                                                                      STREET: 1025 Conr
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 9 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AUDRESSEE: Kenyon & Kenyon
STREET: 1025 Connecticut Avenue, NW - Suite 600
CITY: Washington
STATE. PG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE:
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                                                                                                                                              ADDRESSEB:
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3 IRDDILD 9
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                                                                                                                          B: Kenyon & Kenyon
1025 Connecticut Avenue, NW - Suite 600
                                                                                                                                                                                                                                             KOIKE, Ayumi
OGURA, Kyozo
KOYAMA, Tanetoshi
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SHIMIZU, Naoto
                                                                                                                                                                                                                                                                                                      MURAMATSU, Masayoshi
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RESULT 10
US-08-556-419-14
Sequence 14, Application US/08556419C
Sequence 10, 6093549
; Patent No. 6093549
; GENERAL INFORMATION:
APPLICANT: Ross, Christopher
; APPLICANT: Li, Xiao-Jiang
; APPLICANT: Li, Shi-Hua
                                                                                                                                                                                                                                                                                                               S
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US-09-269-250E-38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 58994
CURRENT APPLICATION NUMBER: US/09/269,250E
CURRENT FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin version 3.1
SEQ ID NO 38
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 5; Conserv
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GENERAL INFORMATION:
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Matches
                    APPLICANT: Sharp, Alan
APPLICANT: Lanahan, Anthony
APPLICANT: Worley, Paul
APPLICANT: Snyder, Solomon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 38,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE:
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FILING DATE: 11-Jun-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 154441/1996
FILING DATE: 14-Jun-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-D
SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
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TELEPHONE: 202-429-1776
TELEFAX: 202-429-0796
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les 4; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: TOPPENETTI, Judith L. REGISTRATION NUMBER: 39,048 REFERENCE/DOCKET NUMBER: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
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INVENTION: Huntingtin-associated protein
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5; Conservative (
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nilarity 57.1%;
Conservative
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100.0%; Pred. No. 4.6e+05;
ive 0; Mismatches 0;
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Pred. No. 4.6e
3; Mismatches
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Sequence 4, Application US/09489760

Patent No. 6878375

GENERAL INFORMATION:
APPLICANT: Rijksuniversiteit Te Leidin
APPLICANT: Hunt, Elsa A.J.M

APPLICANT: Hunt, Donald F

APPLICANT: Heard, Victor H

TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US

CURRENT APPLICATION NUMBER: US/09/489,760

CURRENT FILING DATE: 2000-01-21

EARLIER APPLICATION NUMBER: PCT/NL98/00424

EARLIER FILING DATE: 1998-07-23

NUMBER OF SEQ ID NOS: 16

SOFTWARE: Patentin Ver. 2.1
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US-09-489-760-4
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                                                                                                                                                                                                                                                                    RESULT 12
                                                                                                                                                                                                                                                       US-08-459-568-20
                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 5
                                                                                                                                                                       Sequence 20, Application of Patent No. 5811304
GENERAL INFORMATION:
APPLICANT: Huang, Shi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 52.4%;
Best Local Similarity 80.0%;
Matches 4; Conservative
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LENGTH: 9
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CURRENT APPLICATION UNMBER: US/08/556,419C
CURRENT FILING DATE: 1995-11-09
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: PEPTIDE
LOCATION: (2)..(7)
OTHER INFORMATION: RESIDUES
OTHER INFORMATION: RESIDUES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Histocompatibility antigen FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH:
                                                                                                                      TITLE OF INVENTION: Retir
TITLE OF INVENTION: Zinc
NUMBER OF SEQUENCES: 93
                                                                                                    CORRESPONDENCE ADDRESS:
                               STREET: 4370 La Jo
CITY: San Diego
STATE: California
COUNTRY: U
                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 DDLLE 8
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                                                                                                                                                                                                                                                                                                                                                           1 VLRDDLLEA 9
                                                                                                                                                                                                                                                                                                                                                                                             h 52.4%;
Similarity 55.6%;
5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DDLLQ 5
                                                                                                                                                                                                                                  Application US/08459568
                                                                 E: Campbell and Flores
4370 La Jolla Village Drive, Suite
                                                                                                                                   Retinoblastoma Protein - Interacting Zinc Finger Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Mismatches
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Pred. No. 4.6e+05;
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Pred. No. 4.6e+05;
0; Mismatches 4
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US-08-399-411-20
US-08-399-411-20
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COMPUTER: IDM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/459,568
FILING DATE: 02-UTW-1995
CLASSIFICATION NUMBER: US 08/399,411
APPLICATION NUMBER: DELATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 9-LJ 1264
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20, Application US/08399411
Patent No. 5831008
GENERAL INFORMATION:
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Best Local :
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                                                                                                                                     APPLICATION UNMER: US/08/399,411
FILING DATE: 06-MAR-1995
CLASSIFICATION: 530
ATTORNEY/ACENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: PLJ 1264
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                     TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: :
SEQUENCE CHARACTERISTICS:
LENGTH: 6 aming acids
                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Retinoblastoma Protein -
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U
ZIP: 92122
               TYPE: amino acid TOPOLOGY: linear
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Similarity 80.0%;
4; Conservative
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4370 La Jolla Village Drive, Suite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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Pred. No. 4.6e+05;
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                                                                                                                                                                                                                                                                                                                                                                  Version
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION UNMERS: US/08/516,859A
PILING DATE: 18-AUG-1995
CLASSIFICATION NUMBER: US/08/399,411
PRIOR APPLICATION NUMBER: US/08/399,411
PILING DATE: 06-MAR-1995
PRIOR APPLICATION NUMBER: US/08/399,411
FILING DATE: 06-MAR-1995
PRIOR APPLICATION NUMBER: US/08/292,683
FILING DATE: 18-AUG-1994
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: P.LJ 1776
REFERENCE/DOCKET NUMBER: P.LJ 1776
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPAX: (619) 535-999
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
TYPE: amino acids
                                                                                                                     RESULT 15
US-09-586-472-20
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Best Local Similarity
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US-08-516-859A-20
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US-08-516-859A-20
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                                Sequence 20, Application US/0
Patent No. 6323335
PATENT INFORMATION:
APPLICANT: Huang, Shi
TITLE OF INVENTION: R
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Best Local S
Matches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
ADDRESSEB: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Retinoblastoma Protein - Interacting TITLE OF INVENTION: Zinc Finger Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: C
                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: San Diego
NUMBER OF SEQUENCES: 106
                                                                                                                                                                                                                    4 DDLLE 8
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4; Conserv
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                                                                                                      Application US/09586472
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Conservative
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Pred. No. 4.6e+05;
1; Mismatches 0
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Pred. No. 4.6e+05;
1; Mismatches 0
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Search completed: April 6, 2006, 16:27:37 Job time : 27.6667 secs
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                                                                                                                                                                      Matches
                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                              TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 20:
SEQUENUE CHARACTERISTICS:
LENGTH: 6 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 09/528,706
APPLICATION NUMBER: US 08/516,859
APPLICATION NUMBER: US 08/516,859
FILING DATE: 18-AUG-1995
APPLICATION NUMBER: US 08/399,411
FILING DATE: 06-MAR-1995
APPLICATION NUMBER: US 08/292,683
PILING DATE: 18-AUG-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                    Local Similarity 80.0
                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-LJ 4130
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                       TOPOLOGY: linear SEQ ID NO: 20:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                             4 DDLLE 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: San Diego
STATE: California
                                                                                                                                                                                        50.0%;
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Match Length
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*
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Gapop 10.0 ,
   1867569 seqs, 417829326 residues
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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               US-10-623-176-10
US-10-791-217-5
US-11-007-740-18
US-11-007-740-18
US-10-623-176-46
US-10-623-176-2
US-10-623-176-1
US-11-007-740-20
US-10-623-176-48
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US-10-623-176-49
US-10-623-176-49
US-10-623-176-49
US-10-523-176-49
US-10-523-176-15
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10, Appl
5, Appli
18, Appli
42, Appli
2, Appli
2, Appli
1, Appli
1, Appli
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1, Appli
2, Appli
3, Appli
3, Appli
3, Appli
3, Appli
4, Appli
4, Appli
5, Appli
6, Appli
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	21	21	21	21	21	21	21	21	21	21	21	21	21	22	22	22	23	23
	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	50.0	52.4	52.4	52.4	54.8	54.8
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ALIGNMENTS	US-10-119-536A-109	US-09-833-203-54	US-09-935-430-565	US-09-935-430-219	US-10-700-330-169	US-10-487-268-9	US-10-122-675-6	US-09-989-994-216	US-09-989-994-211	US-09-989-994-208	US-09-990-186-216	US-09-990-186-211	US-09-990-186-208	US-11-007-740-40	US-10-791-217-4	US-10-623-176-14	US-10-623-176-3	US-09-834-765-132
	Sequence 109, App	Sequence 54, Appl	Sequence 565, App	Sequence 219, App	Sequence 169, App	Sequence 9, Appli	Sequence 6, Appli	Sequence 216, App	Sequence 211, App	Sequence 208, App	Sequence 216, App	Sequence 211, App	Sequence 208, App	Sequence 40, Appl	Sequence 4, Appli	Sequence 14, Appl		Sequence 132, App

PRIOR APPLICATION NUMBER: 2003-07-18 PRIOR FILING DATE: 2003-07-18 PRIOR FILING DATE: 2000-01-21 PRIOR FILING DATE: 2000-01-21 PRIOR APPLICATION NUMBER: EP 97202303.0 PRIOR FILING DATE: 1997-07-23 PRIOR APPLICATION NUMBER: PCT/NL98/00424 PRIOR FILING DATE: 1998-07-23 PRIOR APPLICATION NUMBER: J2000-504165 PRIOR FILING DATE: 2000-01-24 NUMBER OF SEQ ID NOS: 77 SOPTWARE: Patentin Ver. 2.1 SEQ ID NO 10 밁 S ; NAME/KEY: SITE ; LOCATION: (1)..(9) US-10-623-176-10 RESULT 1 US-10-623-176-10 Query Match Best Local S Matches 9 Sequence 10, Application US/10623176 Publication No. US20040092446A1 GENERAL INFORMATION: APPLICANT: Goulmy, Els A.J.M. APPLICANT: Hunt, Donald F. APPLICANT: Engelhard, Victor H. TITLE OF INVENTION: HA-1 epitopes and uses thereof FILE REFERENCE: 2183-6047US CURRENT APPLICATION NUMBER: US/10/623,176 CURRENT FILING DATE: 2003-07-18 FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide FEATURE: TYPE: PRT ORGANISM: Artificial Sequence LENGTH: Н 1 VLRDDLLEA 9 Similarity 9; Conserv VLRDDLLEA 9 Conservative 100.0%; <u>,</u> Score 42; Pred. No. 1 Mismatches DB 4; I 1.7e+06; hes 0; Length 0

RESULT 2 US-10-791-217-5

Gaps

0

Sequence 5, Application US/10791217 Publication No. US20040191268A1

GENERAL INFORMATION:

APPLICANT: Goulmy, Elsa A.J.M APPLICANT: Hunt, Donald F APPLICANT: Engelhard, Victor

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RESULT 4

US-10-623-176-42

Sequence 42, Application US/10623176

Publication No. US20040092446A1

GENERAL INFORMATION:

APPLICANT: GOULINY, Els A.J.M.

APPLICANT: Hunt, Donald F.

APPLICANT: Engelhard, Victor H.

TITLE OF INVENTION: HA-1 epitopes and uses the TITLE OF INVENTION: 183-1647US
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Publication No. US20050233350A1
GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1
SEQ ID NO 18
LENGTH: 9
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Best Local
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PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
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CURRENT FILING DATE: 2004-03-02
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TYPE: PRI
TYPE: PRI
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: obtained from KIAA0223 partial complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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9; Conserv
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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Sequence 46, Application US/10623176

Publication No. US20040092446A1

; Publication No. US20040092446A1

; Publication No. US20040092446A1

; Publication No. US20040092446A1

; Publicant INFORMATION:

APPLICANT: Hunt, Donald F.

APPLICANT: Hunt, Donald F.

APPLICANT: Hunt, Donald F.

ITITLE OF INVENTION: HA-1 epitopes and uses thereof

; FILE REFERENCE: 2183-6047US

; CURRENT PILING DATE: 2003-07-18

; PRIOR APPLICATION NUMBER: US/10/623,176

; PRIOR APPLICATION NUMBER: D9/489,760

; PRIOR APPLICATION NUMBER: EP 97202303.0

; PRIOR APPLICATION NUMBER: BP 97202303.0

; PRIOR APPLICATION NUMBER: D97202303.0

; PRIOR APPLICATION NUMBER: D97202303.0
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US-10-623-176-46
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; LOCATION: (1)..(9)
US-10-623-176-42
                                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide; FEATURE: NAME/KEY: SITE LOCATION: (1)..(9)
US-10-623-176-46
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                                                                          Query Match
Best Local S
Matches 8
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SEQ ID NO 42
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PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR PILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
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NUMBER OF SEQ ID NOS: 77
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TYPE: PRT
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OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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8; Conser
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                                                                                              90.5%;
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%; Pred. No. 1.7
0; Mismatches
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Pred. No. 1.7e+06;
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US-10-623-176-2 ; Sequence 2, Application US/10623176 ; Publication No. US20040092446A1

RESULT 6

GENERAL INFORMATION:

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US-10-791-217-2
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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-2
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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: DT 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 1998-07-23
PRIOR FILING DATE: 2000-1-24
                                                                           Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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Best Local Similarity
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SEQ ID NO 2
                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses
                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.1
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TYPE: PRT
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                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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                    VLRDDLLEA 9
VLHDDLLEA 9
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                                                                             Conservative
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                                                                                           88.1%;
88.9%;
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88.9%;
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                                                                         Score 37; DB 4; Length 9;
Pred. No. 1.7e+06;
0; Mismatches 1; Indels
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Pred. No. 1.7e+06;
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Query Match
Best Local Similarity
Thehes 8; Conserve
RESULT 10
US-10-623-176-1
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Sequence 20, Application US/11007740
Publication No. US20050233350A1
GENERAL INFORMATION:
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; SEQ ID NO 20
; LENGTH: 9
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CURRENT FILING DATE: 2004-06-04
PRIOR APPLICATION NUMBER: PCT/NL02/00791
PRIOR FILING DATE: 2002-12-05
PRIOR APPLICATION NUMBER: EP 01204704.9
PRIOR FILING DATE: 2001-12-05
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Goulmy, Elsa A.J.M.
APPLICANT: Goulmy, Elsa A.J.M.
TITLE OF INVENTION: Minor histocompatibility antigen HA-1: target antigen for immunc
TITLE OF INVENTION: of tumors
FILE REFERENCE: 2183-6479US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Goulmy, E18a
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
NUMBER OF SEQ ID NOS: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: Exon
                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                              TYPE: PRT
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                                                                                                                                                                         Score 37;
Pred. No.
                                                                                                                                                         Mismatches
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1.7e+06;
                                                                                                                                                                                           Length 9;
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Sequence 1, Application US/10623176
Publication No. US20040092446A1
GENERAL INFORMATION:

APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.

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CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 2000-01-21
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR FILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: JP 2000-504165
PRIOR FILING DATE: 2000-01-24
NUMBER OF SEQ ID NOS: 77
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Best Local Similarity
Matches 8; Conser
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Publication No. US20040191268A1
GENERAL INFORMATION:
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SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 1
                                                                                           Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Goulmy, Elsa A.J.M
APPLICANT: Hunt, Donald F
APPLICANT: Engelhard, Victor H
TITLE OF INVENTION: The HA-1 Antigen
FILE REFERENCE: 2183-4285US
                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/10/791,217
CURRENT FILING DATE: 2004-03-02
PRIOR APPLICATION NUMBER: US/09/489,760
PRIOR FILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: HA-1 epitopes and uses thereof FILE REFERENCE: 2183-6047US
                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR FILING DATE: 1998-07-23
NUMBER OF SEQ ID NOS: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide OTHER INFORMATION: wherein X can be R or H
                                                                                                                                                           PEATURE:
NAME/KBY: MISC FEATURE
LOCATION: (3)...(3)
OTHER INFORMATION: Xaa is HISTIDINE OR ARGININE RESIDUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: SITE
LOCATION: (1)...
                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial sequence
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                                                                                                                                                                                                                                                   OTHER INFORMATION: obtained from histocompatibility antigen
                                                                                                                                                                                                                                                                       FEATURE:
                                                                       Local Similarity
les 8; Conserv
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                                     1 VLRDDLLEA 9
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88.9%;
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Pred. No. 1.7e+06;
                                                                             0
                                                                       Score 36; DB 4;
Pred. No. 1.7e+06;
0; Mismatches 1
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; FEATURE:
; NAME/KEY: MISC FEATURE
; COCATION: (3)...(3)
; OTHER INFORMATION: Xaa represents a histidine (H)
US-11-007-740-29
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APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epicopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR FILING DATE: 1900-01-21
PRIOR PILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EPT/NL98/00424
PRIOR FILING DATE: 1998-07-23
PRIOR APPLICATION NUMBER: PCT/NL98/00424
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 1998-07-23
PRIOR PILING DATE: 2000-01-24
NUMBER: DE 2000-01-24
NUMBER: DE 2000-01-24
NUMBER: DE 2000-01-24
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US-11-007-740-29
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                                                                                                                                    ; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-48
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Best Local Similarity
Matches 8; Conserv
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                                                           Best Loc
Matches
                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 48, Application US/10623176
Publication No. US20040092446A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goulmy, E18a
TITLE OF INVENTION: METHOD FOR TYPING OF MINOR HISTOCOMPATIBILITY ANTIGEN HA-1
FILE REFERENCE: 2799/58994-A
CURRENT APPLICATION NUMBER: US/11/007,740
CURRENT FILING DATE: 2004-12-08
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR APPLICATION NUMBER: 09/269,250
PRIOR FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 40
                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
                                                                                                                                                                                               OTHER INFORMATION: FEATURE:
                                                                                                                                                                                                                                    TYPE: PRI
TYPE: PRI
ORGANISM: Artificial Sequence
FEATURE:
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les 7; Conserv
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3 RDDLLEA 9
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                                                           Conservative
                                                                                                                                                                                                                   Description of Artificial Sequence: HA-1 peptide
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88.9%;
                                                       81.0%; 5u
100.0%; Pr
                                                             score 34; DB; Pred. No. 1.7
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Pred. No. 1.7e+06;
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                                                                                                     DB 4;
                                                                                 .7e+06;
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US-10-623-176-23

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APPLICANT: HUNTER, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
PILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR PILING DATE: 1997-07-23
PRIOR APPLICATION NUMBER: EP 97202303.0
PRIOR APPLICATION NUMBER: DE 97202303.0
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; NAME/KEY: SITE
; LOCATION: (1)..(9)
US-10-623-176-23
                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 41
LENGTH: 9
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Publication No. US20040092446A1
GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 7; Conservative
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Publication No. US20040092446A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Goulmy, Els A.J.M.
APPLICANT: Hunt, Donald F.
APPLICANT: Engelhard, Victor H.
TITLE OF INVENTION: HA-1 epitopes and uses thereof
FILE REFERENCE: 2183-6047US
CURRENT APPLICATION NUMBER: US/10/623,176
CURRENT FILING DATE: 2003-07-18
PRIOR APPLICATION NUMBER: 09/489,760
PRIOR PILING DATE: 2000-01-21
PRIOR FILING DATE: 2000-01-21
                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: EP 97202303.0 PRIOR FILING DATE: 1997-07-23 PRIOR APPLICATION NUMBER: PCT/NL98/00424 PRIOR FILING DATE: 1998-07-23 PRIOR APPLICATION NUMBER: JP 2000-504165 PRIOR APPLICATION NUMBER: JP 2000-504165 PRIOR FILING DATE: 2000-01-24 NUMBER OF SEQ ID NOS: 77
                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: HA-1 peptide
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100.0%; Pred. No. 1.7e+06;
7ative 0; Mismatches 0;
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Search completed: April 6, 2006, 16:50:29 Job time: 90.3333 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             on:
      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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      Published_Applications_AA_New:*

1: /SIDSS/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

2: /SIDSS/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /SIDSS/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

4: /SIDSS/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

5: /SIDSS/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

6: /SIDSS/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

7: /SIDSS/ptodata/2/pubpaa/US11_NEW_PUB.pep:*

8: /SIDSS/ptodata/2/pubpaa/US60_NEW_PUB.pep:*
    BLOSUM62
Gapop 10.0 , Gapext 0.5
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42
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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      9888777755555555777777499
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US-11-010-748A-12
US-11-010-748A-11
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US-11-225-686-208
US-11-225-686-211
US-11-225-686-211
US-11-225-099-211
US-11-225-099-211
US-11-225-099-216
US-11-229-099-216
US-11-229-099-167A-219
US-11-129-143-168
US-11-129-143-173
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Sequence 12, Appl
Sequence 11, Appl
Sequence 210, Appl
Sequence 208, Appl
Sequence 2116, App
Sequence 216, App
Sequence 2116, App
Sequence 565, App
Sequence 168, App
Sequence 170, App
Sequence 171, App
Sequence 173, App
Sequence 173, App
Sequence 173, App
Sequence 174, App
Sequence 184, App
Sequence 65, App1
Sequence 66, App1
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RESULT 2
US-11-010-748A-11
; Sequence 11, Application US/11010748A
; Publication No. US20050244421A1
; GENERAL INFORMATION:
; APPLICANT: Merck Patent GmbH
; APPLICANT: MOLL, Heidrun
; APPLICANT: MOLL, Heidrun
; APPLICANT: SCHARM, Burkhard

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Query Match Best Local Matches Qy 1 Db 1	ORGANI FEATUR FEATUR NAME/K LOCATI OTHER OTHER OTHER	APPLICANT: APPLICANT: APPLICANT: APPLICANT: FIFLE OF II FIFLE OF II FILE REFERI CURRENT APPLICANT: PRIOR APPLICANT: PRIOR APPLICANT: PRIOR APPLICANT PRIOR FILII FRIOR FILII F	ESULT 1 (S-11-010- Sequence Publicat GENERAL	22222666666666666666666666666666666666
ch Similarity 9; Conservat 9; Conservat 1 VLRDDLLEA 9 1 VLRDDLLEA 9	Homo sapi MISC_FEAT (1)(9) ORMATION: ORMATION: A-12	: MELCA FOL : MOLLA, He : MOLLA, He : MOLLA, He : MOLLA, HE : SCHARM, INVENTION: RENCE: MER- PPLICATION NU ILICATION NU LICATION NU LICATION NU LICATION U LICATION U	() * 29	119 4 5 5 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
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%; Score 42; DB 7; Length 9)%; Pred. No. 1.4e+05; 0; Mismatches 0; Indels	minor histocompatibility antigens pitopes	idrun iTTER, Wolfgang idrun Burkhard COMPOSITIONS AND METHODS FOR MODULATING 136 COMPOSITIONS AND METHODS FOR MODULATING 137 COMPOSITIONS AND METHODS FOR MODULATING 138 COMPOSITIONS AND METHODS FOR	ALIGNMENTS 1010748A	US-10-927-634-63 US-11-225-686-2954 US-11-225-686-3917 US-11-225-686-3918 US-11-225-686-3918 US-11-225-686-3918 US-11-225-686-3932 US-11-225-686-3941 US-11-225-686-3941 US-11-202-009-3914 US-11-202-009-3914 US-11-202-009-3916 US-11-202-009-3916 US-11-202-009-3918 US-11-202-009-3918 US-11-202-009-3919 US-11-202-009-3919 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-202-009-3941 US-11-045-024-3698 US-11-045-024-3698 US-11-045-024-3698 US-11-045-024-5716
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Gaps	characterized	NE RESPONSE		53, Ap 2854, 3877, 3914, 3915, 3916, 3932, 3941, 2954, 3914, 3915, 3916,
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RESULT, 4
US-11-096-706-52
; Sequence 52, Application US/11096706
; Publication No. US20050245476A1
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Matches
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                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: A-890A
CURRENT APPLICATION NUMBER: US/11/019,027
CURRENT FILING DATE: 2004-12-21
PRIOR APPLICATION NUMBER: 60/605,902
PRIOR FILING DATE: 2004-08-31
PRIOR APPLICATION NUMBER: 60/531,714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: ZHOU, Hongxing
APPLICANT: ZHOU, Chen
APPLICANT: COSMAN, David J.
APPLICANT: CORTER, Paul
APPLICANT: CARTER, Paul
APPLICANT: MARTIN, Francis H.
APPLICANT: MARTIN, FRANCIS H.
TITLE OF INVENTION: METHODS OF IDENTIFYING FUNCTIONAL ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: YAN, APPLICANT: SHE
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                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 2003-
NUMBER OF SEQ ID NOS: 73
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OTHER INFORMATION: Human minor histocompatibility antigens HA-1 characterized by T
OTHER INFORMATION: ell epitopes
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ORGANISM: Homo sapiens
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ORGANISM: Artificial sequence
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88.9%;
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100.0%; Pred. No. 1.4e+05;
tive 0; Mismatches 0;
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US-11-225-686-208
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US-11-225-686-211
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APPLICANT: Sangamo Biosciences, Inc.
APPLICANT: Sangamo Biosciences, Inc.
APPLICANT: Collingwood, Trevor
IITLE OF INVENTION: Treatment of Neuropathic Pain with Zinc Finger Proteins
FILE REFERENCE: 019496-008220US
CURRENT APPLICATION NUMBER: US/11/096,706
CURRENT FILING DATE: 2005-04-01
PRIOR APPLICATION NUMBER: US 60/560,535
PRIOR APPLICATION NUMBER: US 60/560,535
PRIOR APPLICATION NUMBER: US 60/576,757
PRIOR APPLICATION NUMBER: US 60/576,757
PRIOR PILING DATE: 2004-06-02
NUMBER OF SEQ ID NOS: 227
RUMBER OF SEQ ID NOS: 227
NUMBER OF SEQ ID NOS: 227
NUMBER OF SEQ ID NOS: 227
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Publication No. US20060019349A1

GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/225,686
CURRENT FILING DATE: 2001-11-20
PRIOR PPLICATION UNMER: US/09/98,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
CORTWANDE: DESCRIPT US: 200
Sequence 211, Application US/11225686

Publication No. US20060019349A1

GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/225,686
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SEQ ID NO 208
LENGTH: 7
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TYPE: PRT
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Pred. No. 1.4e+05;
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RESULT 8
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                                      APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/202,009
CURRENT FILING DATE: 2005-08-11
PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/225,686
CURRENT FILING DATE: 2005-09-12
PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 216
                                                                                                                                                                                                                                                                          Sequence 208, Application US/11202009 Publication No. US20060024726A1 GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US/09/989,994
PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 211
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Best Local Similarity
Matches 4; Conser
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Best Local Similarity
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: example ZFP
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ORGANISM: Artificial Sequence
FEATURE:
ORGANISM: Artificial Sequence
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Pred. No. 1.4e+05;
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Pred. No. 1.4e+05;
1; Mismatches 1
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                                                                                                   ; OTHER INFORMATION: Description of Artificial Sequence: example US-11-202-009-216
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US-11-202-009-216
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PRIOR FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 211
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Publication No. US20060024726A1
GENERAL INFORMATION:
APPLICANT: LIU, Olang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLECTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: B325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/11/202,009
CURRENT FILING DATE: 2005-08-11
                                Query Match
Best Local Similarity
Matches 4; Conserv
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APPLICANT: LIU, Qia
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Publication No. US20060024726A1
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
PRATURE:
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TYPE: PRT
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3 RDDLLE 8
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                                  Conservative
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                                Score 21; DB 7;
Pred. No. 1.4e+05;
1; Mismatches 1
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Pred. No.
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Pred. No. 1.4e+05;
1; Mismatches 1
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                                                                  DB 7;
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                                  1; Indels
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PRIOR FILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR APPLICATION NUMBER: 60/282,739
PRIOR APPLICATION NUMBER: 60/282,739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 219
LENGTH: 9
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Publication No. US20060018917A1
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 565, Applic Publication No. US20 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                           APPLICANT:
APPLICANT:
APPLICANT: CHALLITA-EID, PIA
APPLICANT: JAKOBOVITZ, AYA
APPLICANT: JAKOBOVITZ, AYA
TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 511582005004
CURRENT FOR APPLICATION NUMBER: US/10/989,767A
CURRENT FILING DATE: 2004-11-15
PRIOR APPLICATION NUMBER: 10/277,292
PRIOR PILING DATE: 2002-10-21
PRIOR PILING DATE: 2002-10-21
PRIOR APPLICATION NUMBER: 09/935,430
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
PRIOR PILING DATE: 2001-08-22
PRIOR PILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: 60/227,098
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TITLE OF INVENTION: NUCLEIC ACID AND CORRESPONDING PROTEIN NAMED 158P1D7
TITLE OF INVENTION: USEFUL IN THE TREATMENT AND DETECTION OF BLADDER AND
TITLE OF INVENTION: OTHER CANCERS
FILE REFERENCE: 511582005004
CURRENT PEPLICATION NUMBER: US/10/989,767A
CURRENT FILING DATE: 2004-11-15
CURRENT FILING DATE: 2004-11-15
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                                                                                                                                                                                                                                                                                                                                                              APPLICANT: FARIS, MARY
APPLICANT: HUBERT, REN
APPLICANT: RAITANO, AF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 LMKSDLVE 8
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RAITANO, ARTHUR
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LEVIN, ELANA
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RAITANO, ARTHUR
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Pred. No. 1.4e+05;
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PRIOR APPLICATION NUMBER: 60/282,739
PRIOR FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 700
SOPTWARE: Patentin Ver. 2.1
SEQ ID NO 565
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                            CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
SOPTWARE: PatentIn version 3.1
SEQ ID NO 169
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Publication No. US20050266518A1
GENERAL INFORMATION:
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SEQ ID NO 168
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Best Local Similarity 37.1
Matches 3; Conservative
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Best Local S
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APPLICANT:
APPLICANT:
                                                                                                                          APPLICANT: BERRY, Alan
APPLICANT: BRETZEL, Weriner
APPLICANT: HUMBELIN, Markus
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: LOPEZ-ULIBARRI, Rual
APPLICANT: MAYER, Anne F.
APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/11/129,143
CURRENT FILING DATE: 2005-05-13
NUMBER OF SEQ ID NOS: 197
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APPLICANT: BRETZEL, Werner
APPLICANT: HUMBELIN, Markus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: YELISEEV, Alexei A.
TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION
FILE REFERENCE: C38435/121966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 5
TYPE: PRT
ORGANISM: Bradyrhizobium japonicum
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
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o. US20050266518A1
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60.0%; Pred. No. 1.4e+05;
tive 2; Mismatches 0
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Pred. No. 1.4e
4; Mismatches
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US-11-129-143-170

Sequence 170, Application US/11129143

Publication No. US20050266518A1

GENERAL INFORMATION:

APPLICANT: BERTY, Alan

APPLICANT: HUMBELLN, Markus

APPLICANT: HUMBELLN, Markus

APPLICANT: LOPEZ-ULIBARRI, Rual

APPLICANT: YELISETY, Alexei A.

TITLE OF INVENTION: IMPROVED ISOPRENOID PRODUCTION

FILE REFERENCE: C38435/121966

CURRENT APPLICATION NUMBER: US/11/129,143

CURRENT FILING DATE: 2005-05-13

NUMBER OF SEQ ID NOS: 197

SOFTWARE: Patentin version 3.1

LENGTH: 5

TYPE: PRT

ORGANISM: Bacillus stearothermophilus

US-11-129-143-170
Search completed: April 6, 2006, 16:51:12 Job time: 12.333 secs
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US-11-129-143-169
                                                                                                                                                             Query Match 47.6%; Score 20; DB 7; Length 5; Best Local Similarity 60.0%; Pred. No. 1.4e+05; Matches 3; Conservative 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 47.6%; Score 20; DB 7; Length 5; Best Local Similarity 60.0%; Pred. No. 1.4e+05; Matches 3; Conservative 2; Mismatches 0; Indels
                                                                               4 DDLLE 8
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1 DDILD 5
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1 DDILD 5
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